



Fig. 1

09022860 64E22860

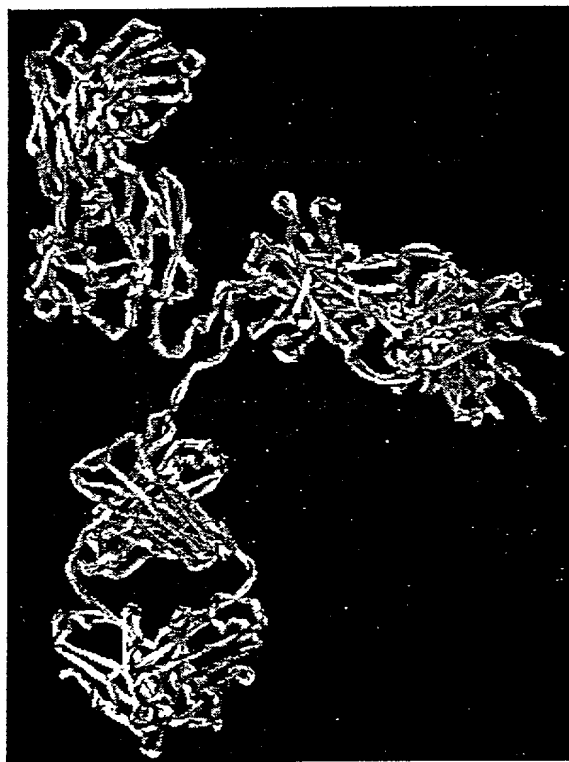


Fig. 2

K2 Fragment:

| | XmaI | | Modified XmaI |
|----|--------|---|---------------|
| 5' | CCGGGC | AGA AGG GCA AGT CTG CAT AGA AGG GCA AGT ATG AAG GCA | 3' |
| 3' | | CG TCT TCC CGT TCA GAC GTA TCT TCC GCT TCA TAC TTC CGTGGCC | 5' |
| | | <u>Arg Arg Ala Ser</u> Leu His <u>Arg Arg Ala Ser</u> Met Lys Ala | |

Fig. 3

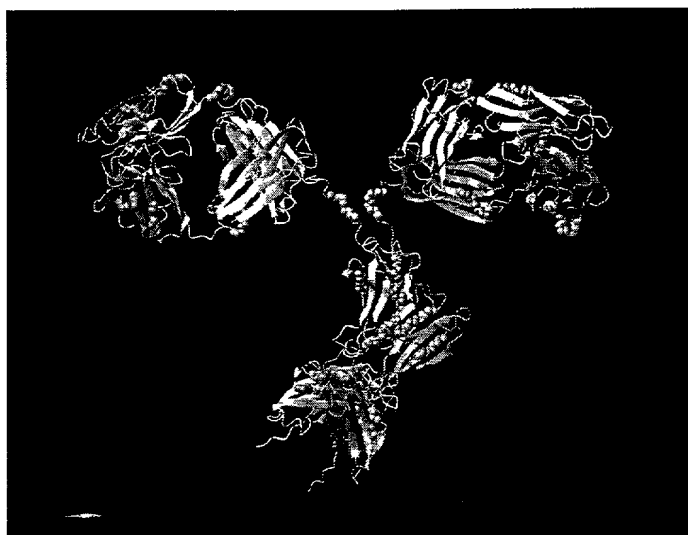


Fig. 4

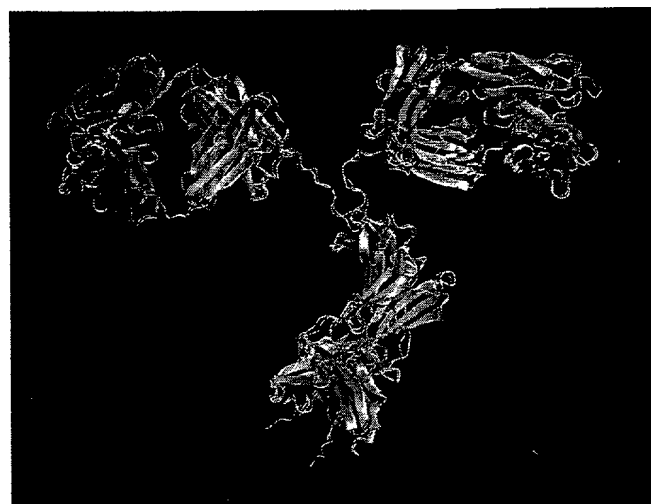


Fig. 5

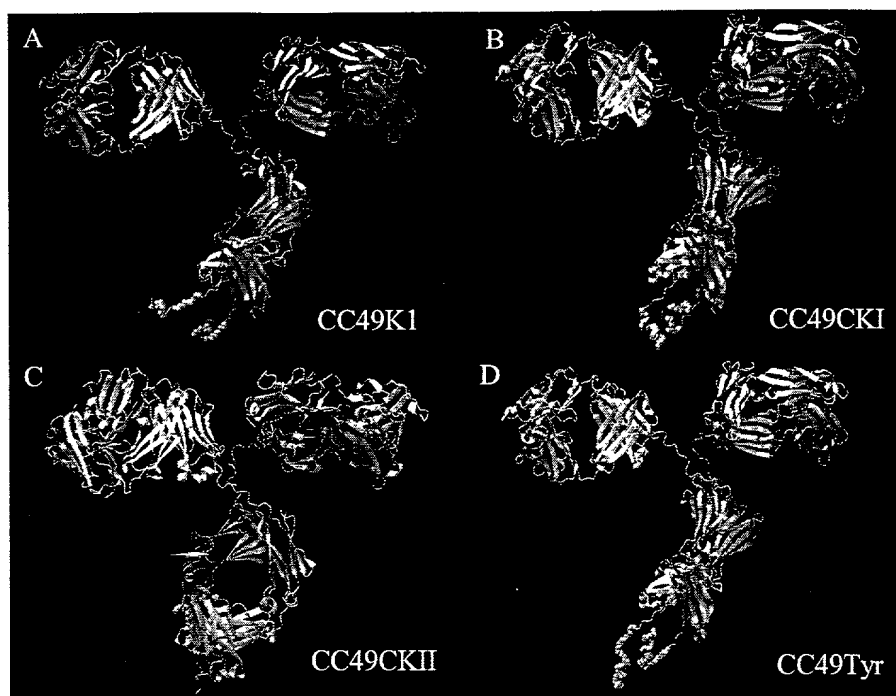


Fig. 6



Fig. 7

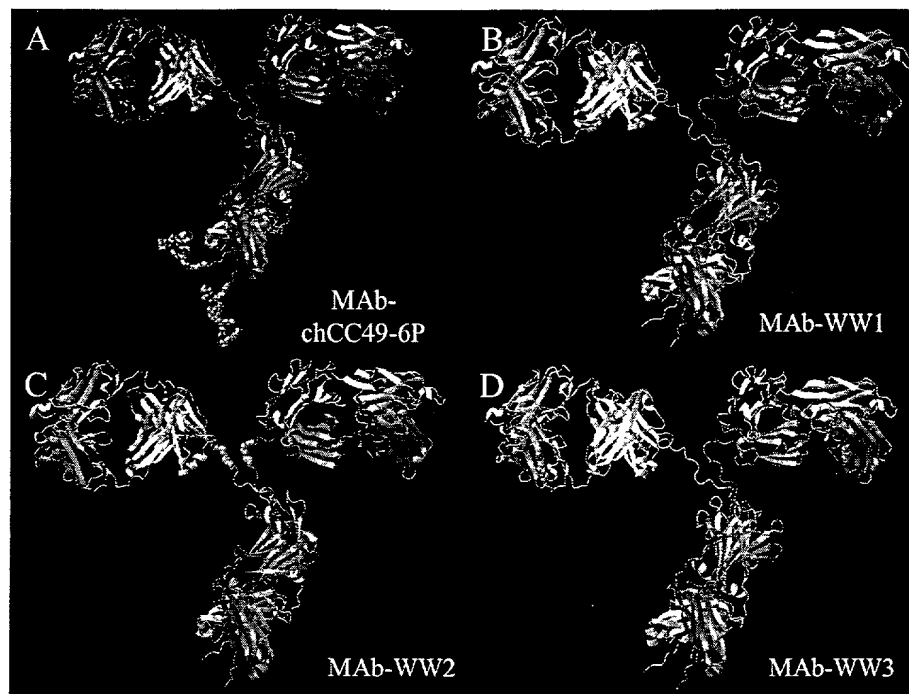


Fig. 7A-D

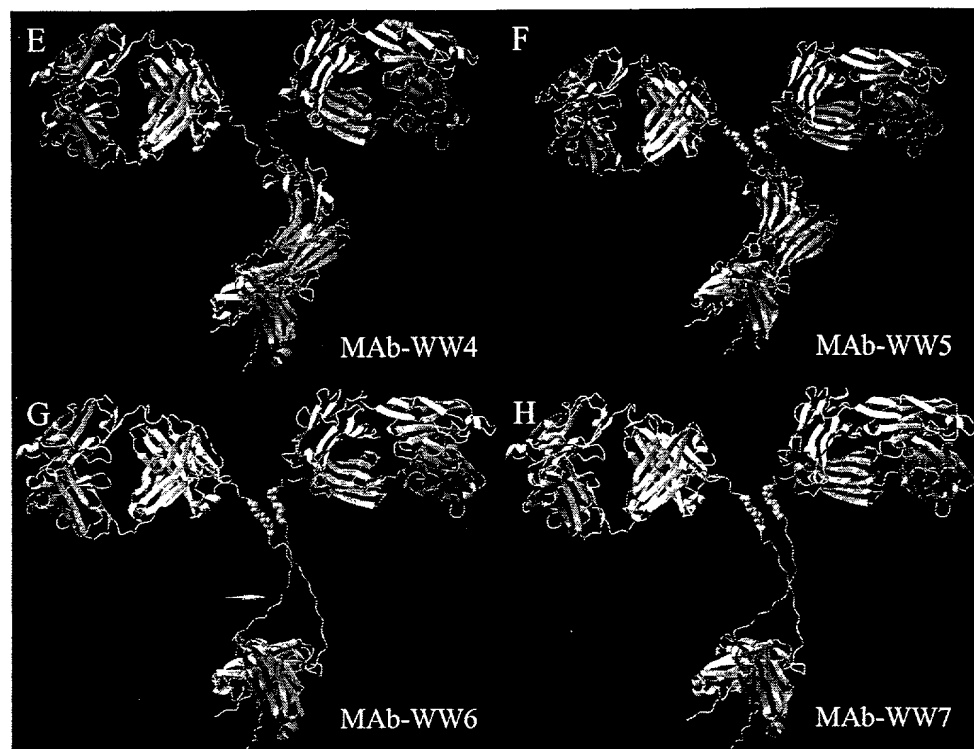


Fig. 7E-H

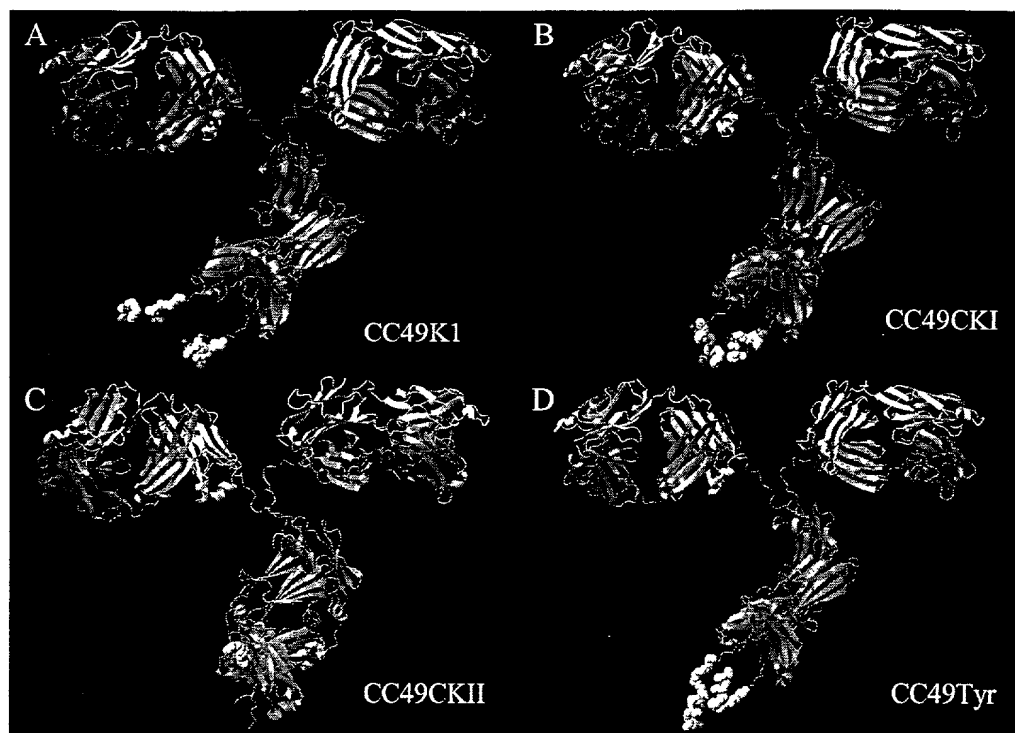


Fig. 8



Fig. 9

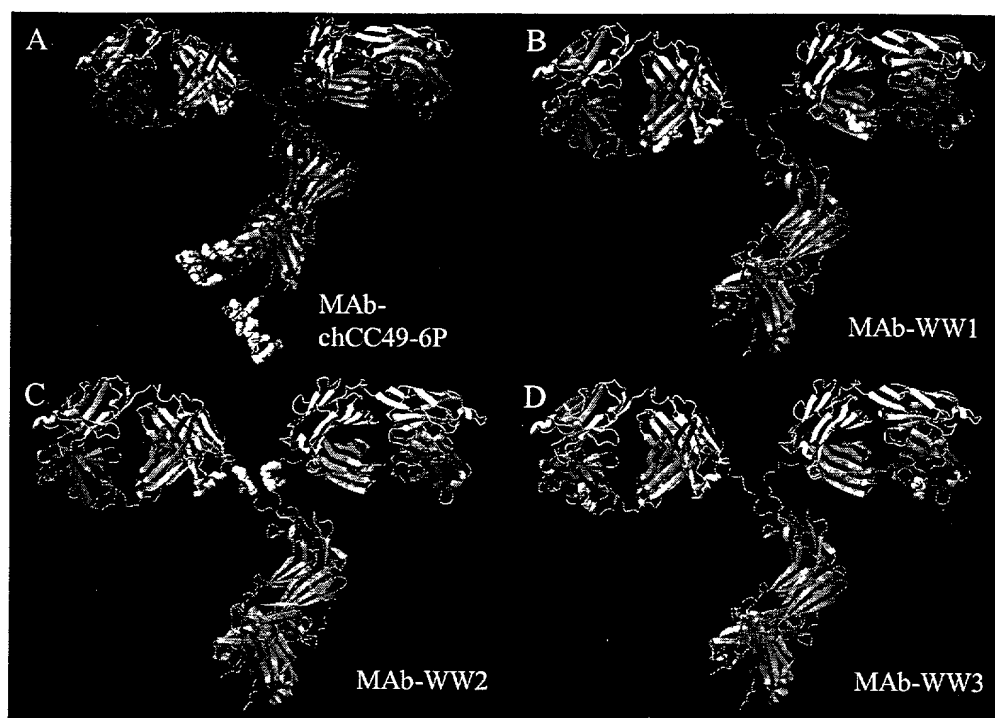


Fig. 9A-D

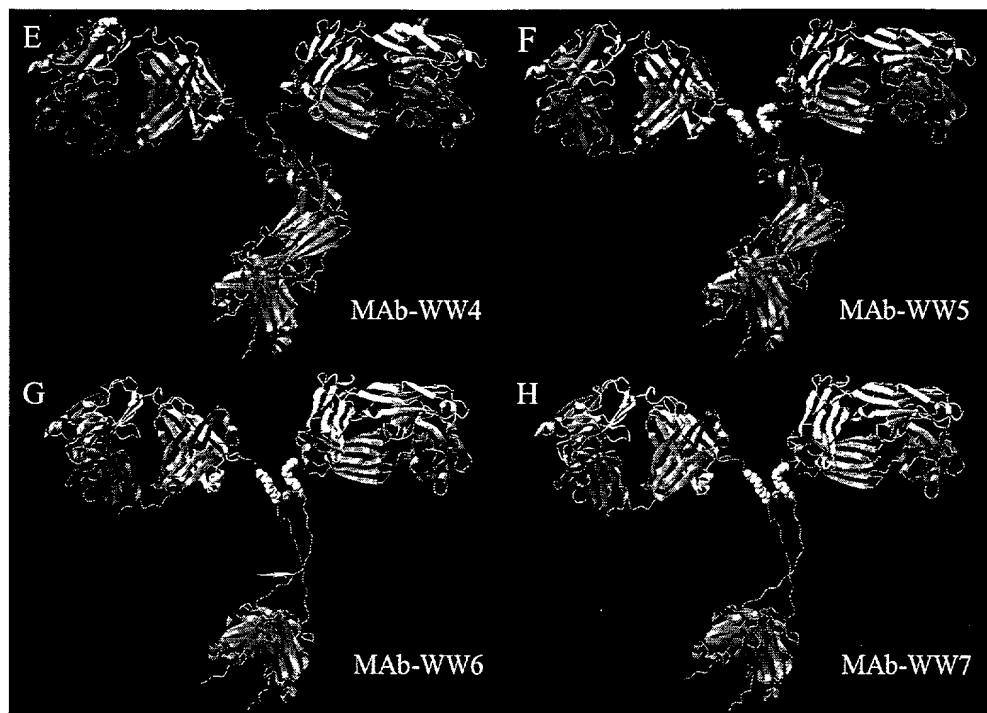


Fig. 9E-H

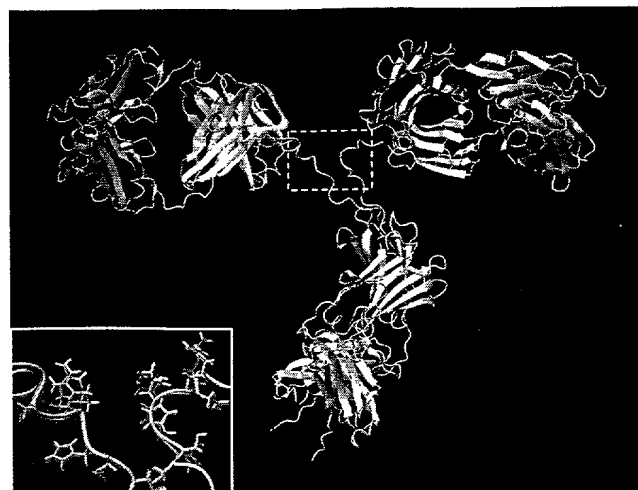


Fig. 10

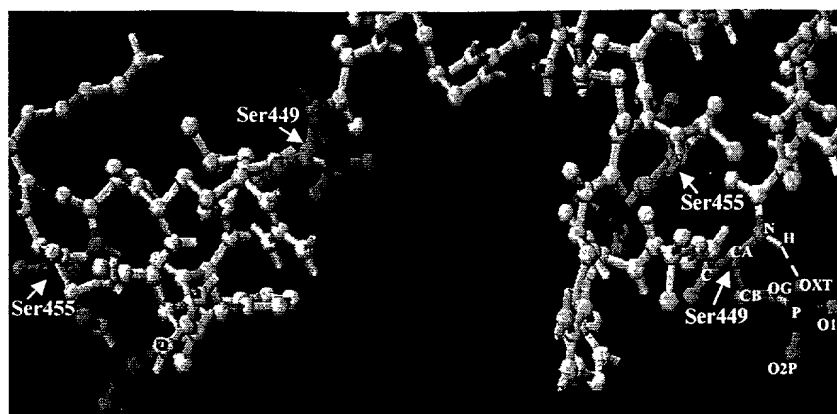
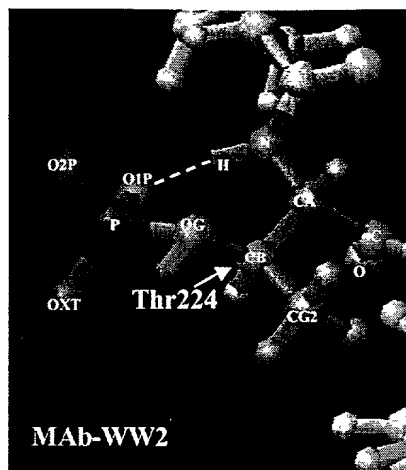


Fig. 11

A



B

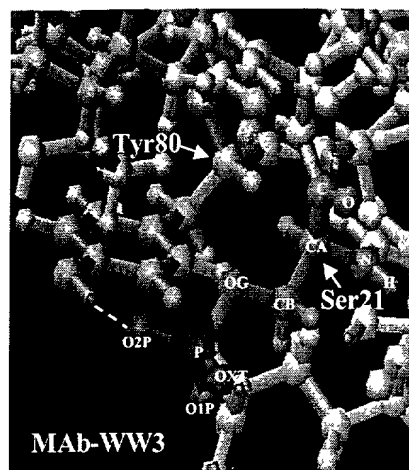


Fig. 12

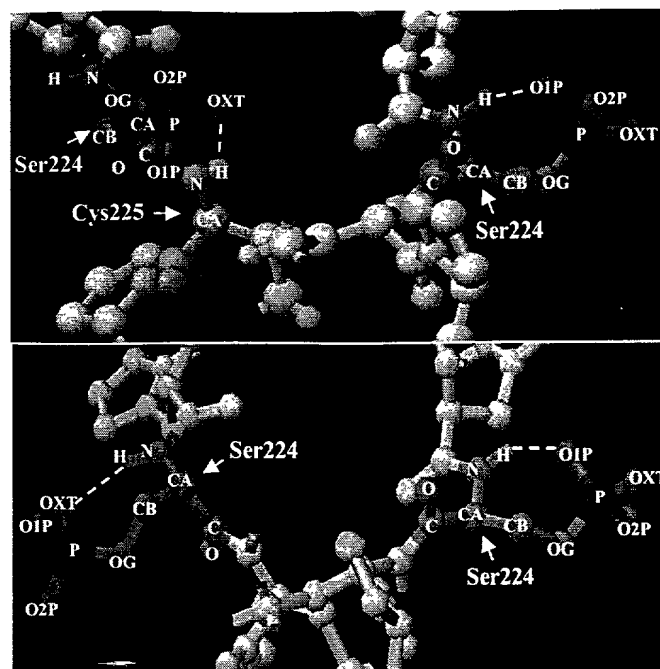
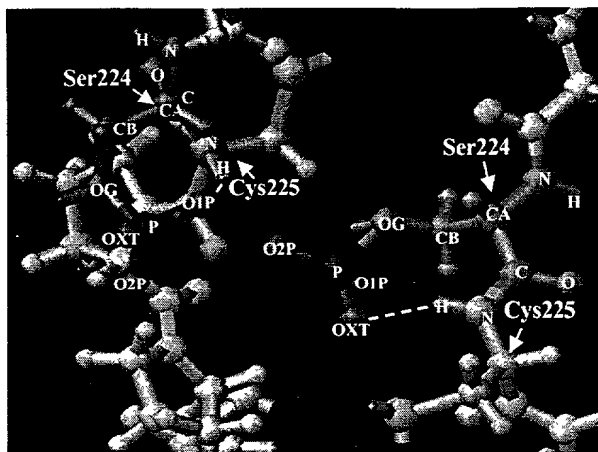


Fig. 13

A



B

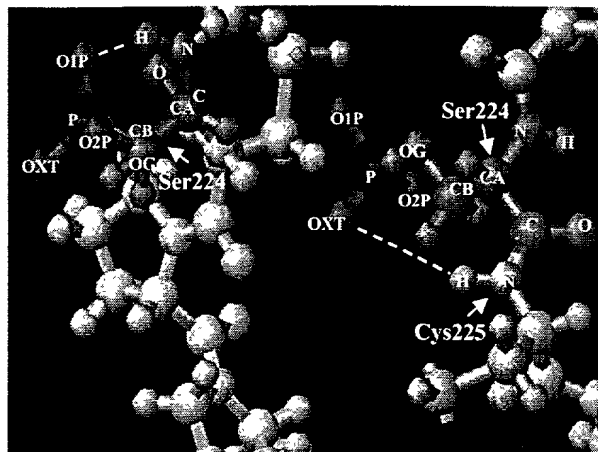
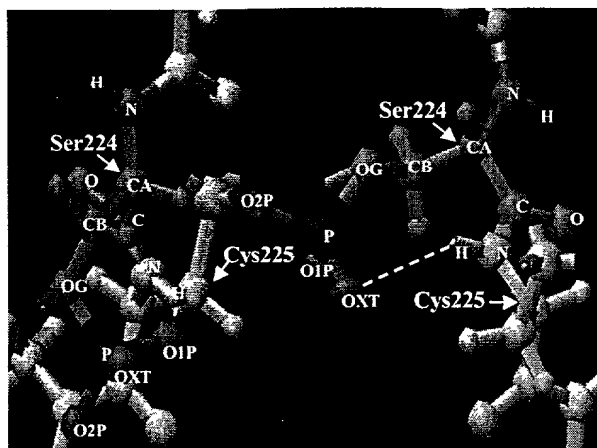


Fig. 14

A



B

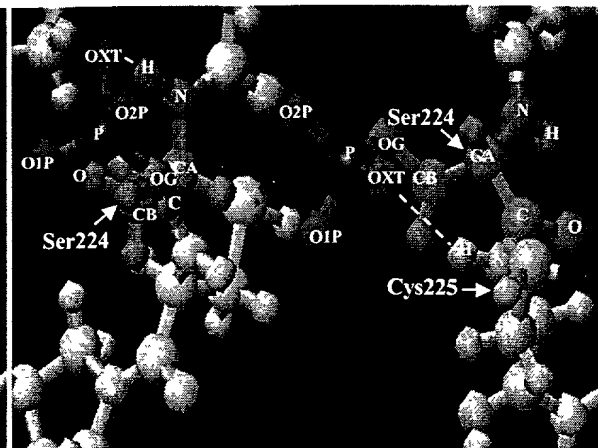


Fig. 15

B

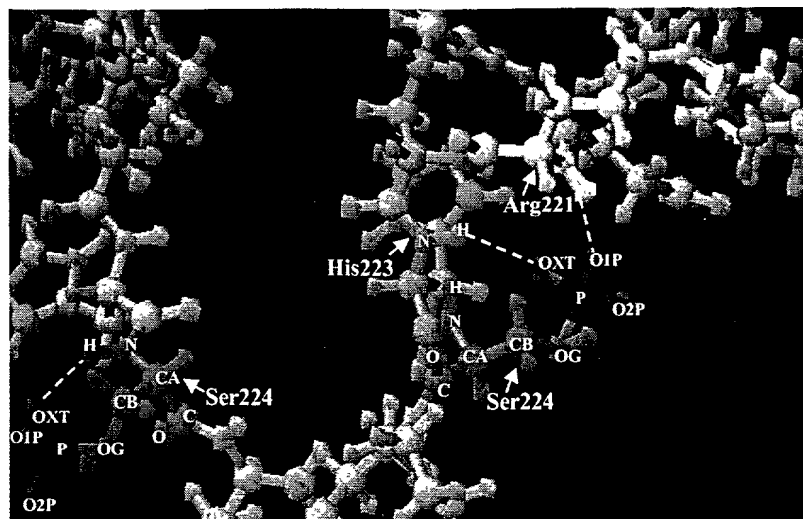


Fig. 16

A

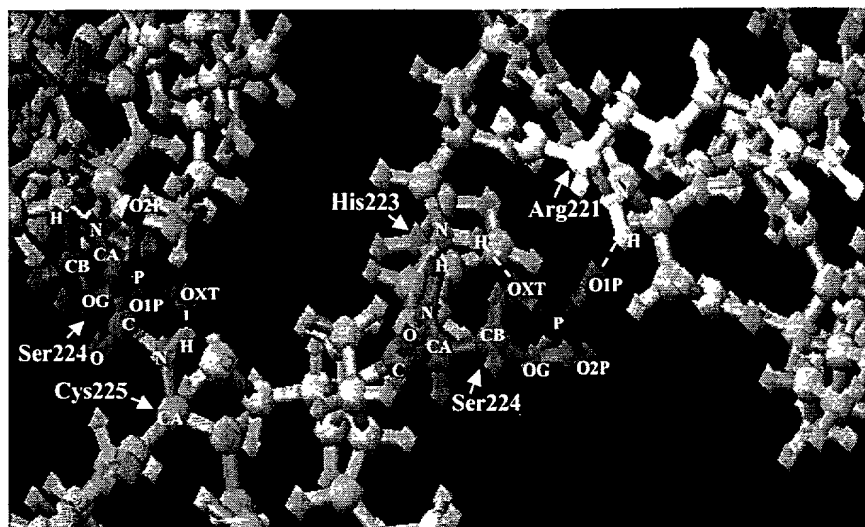


Fig. 16A

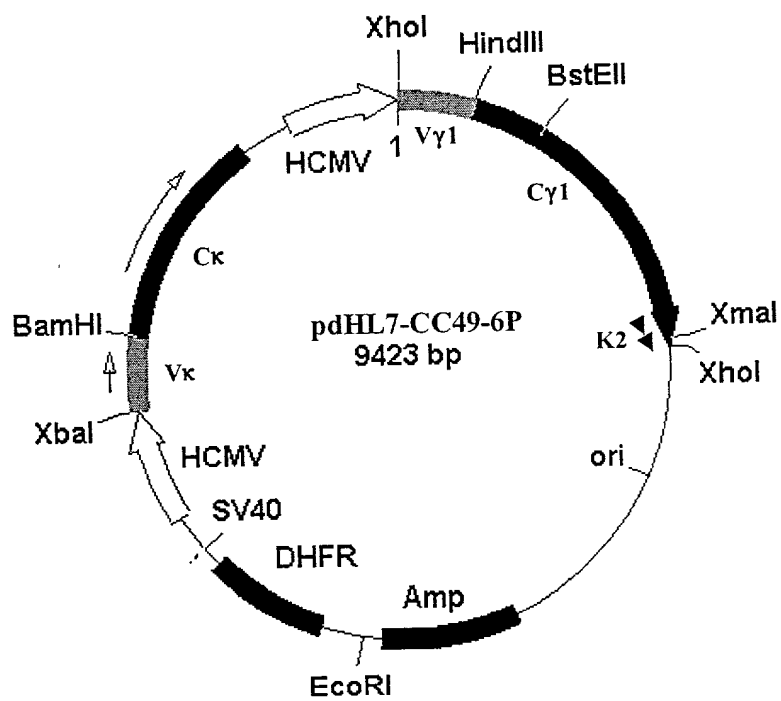


Fig. 17

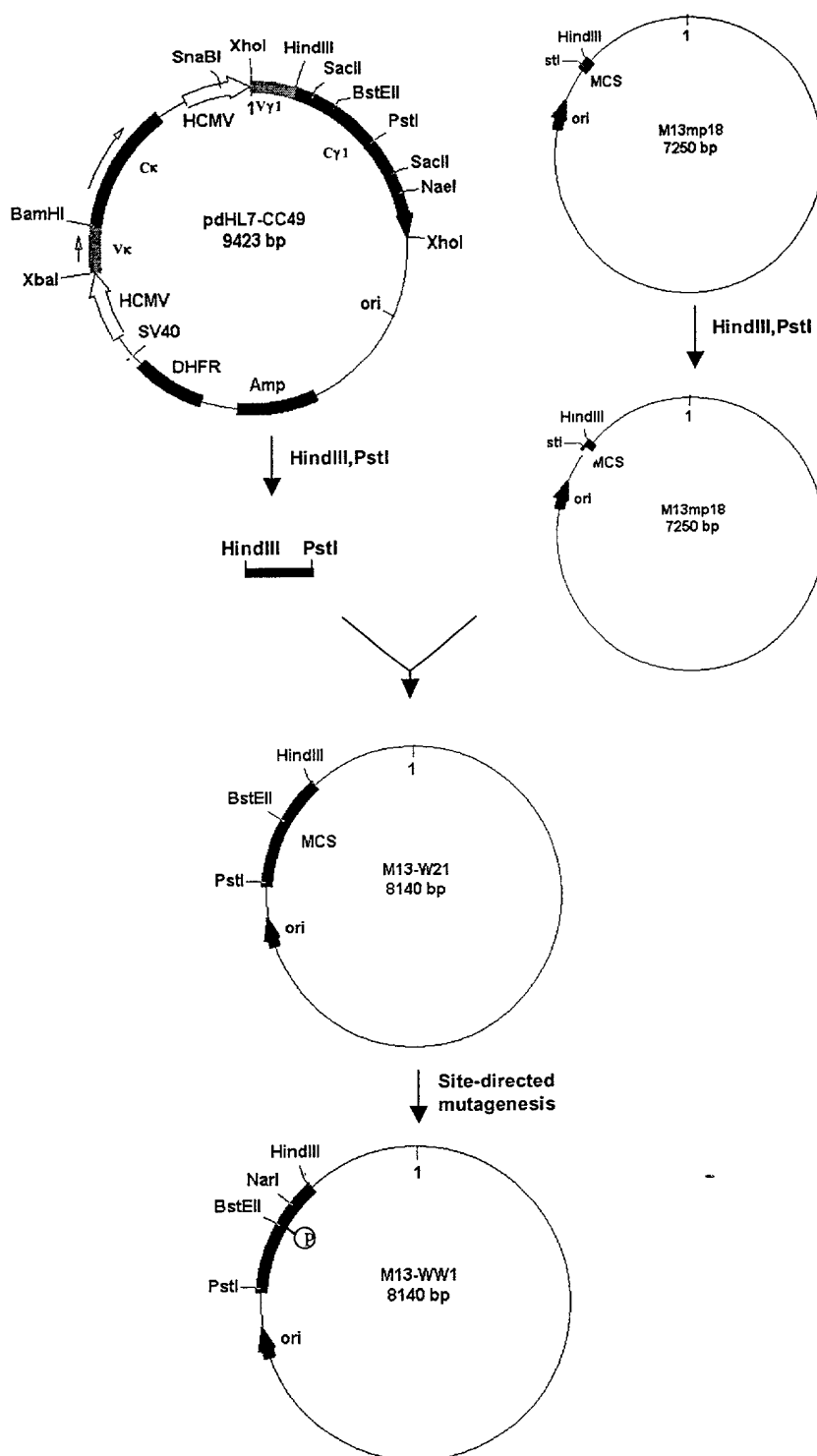


Fig. 18A

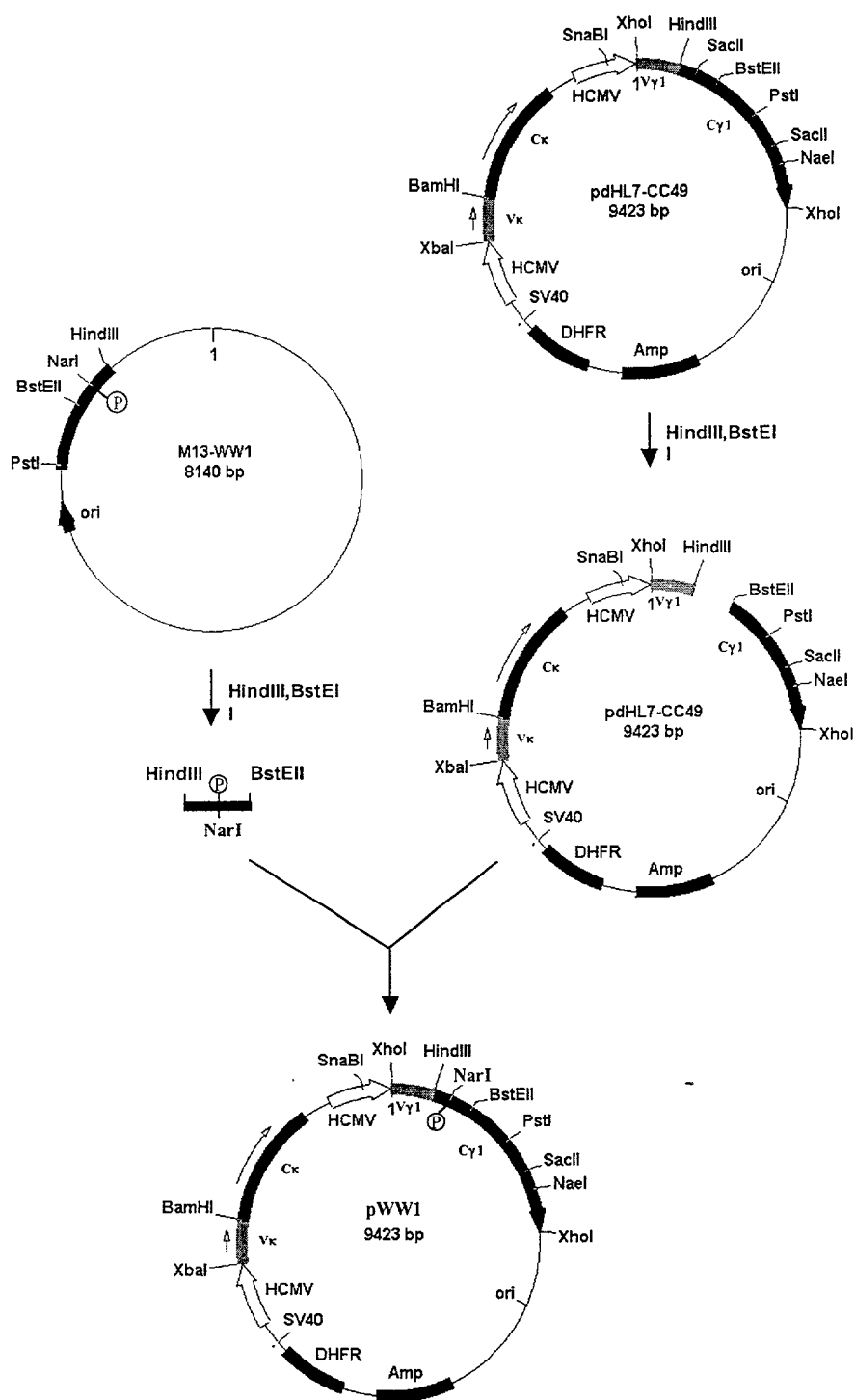


Fig. 18B

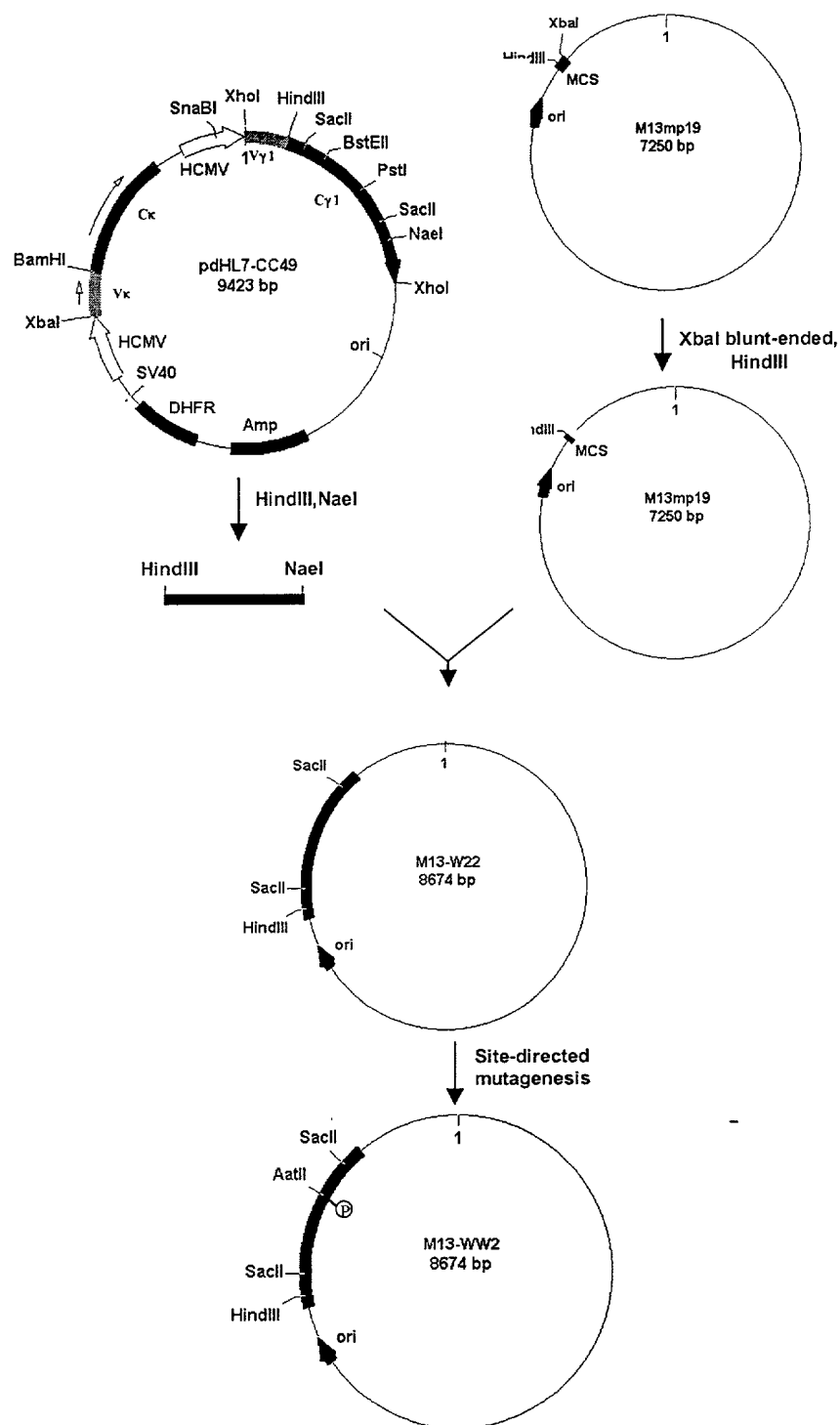


Fig. 19A

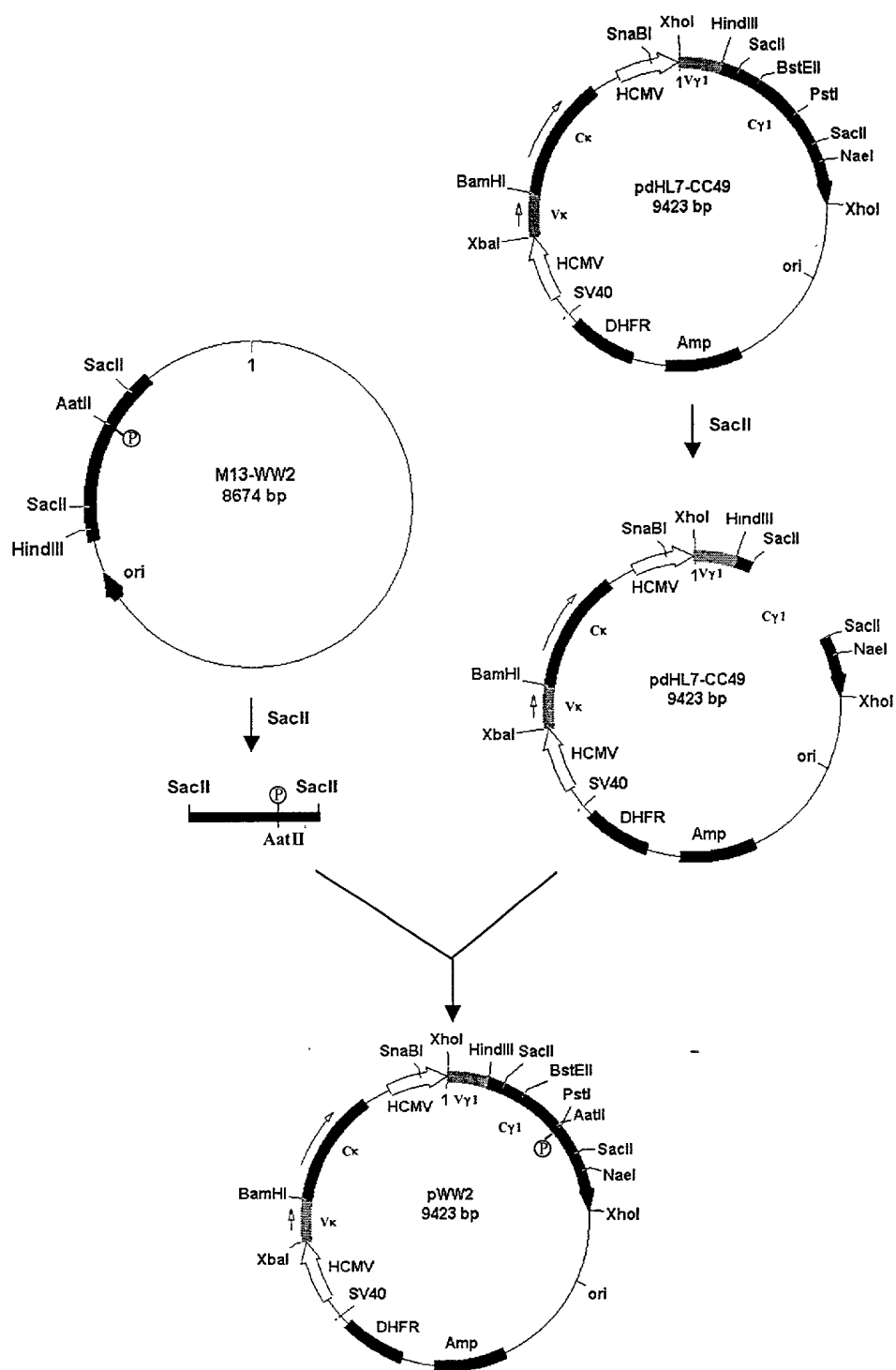


Fig. 19B

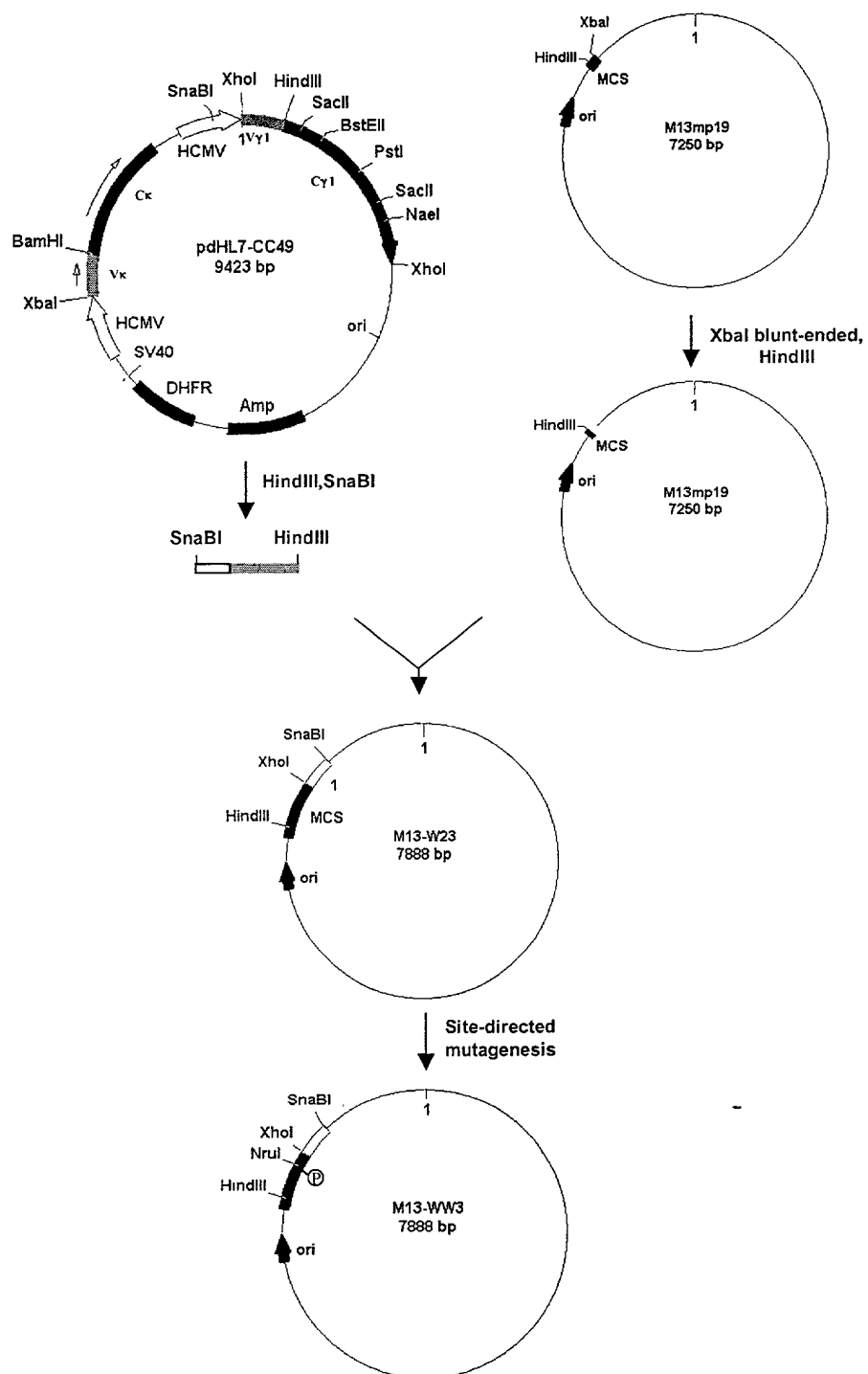


Fig. 20A

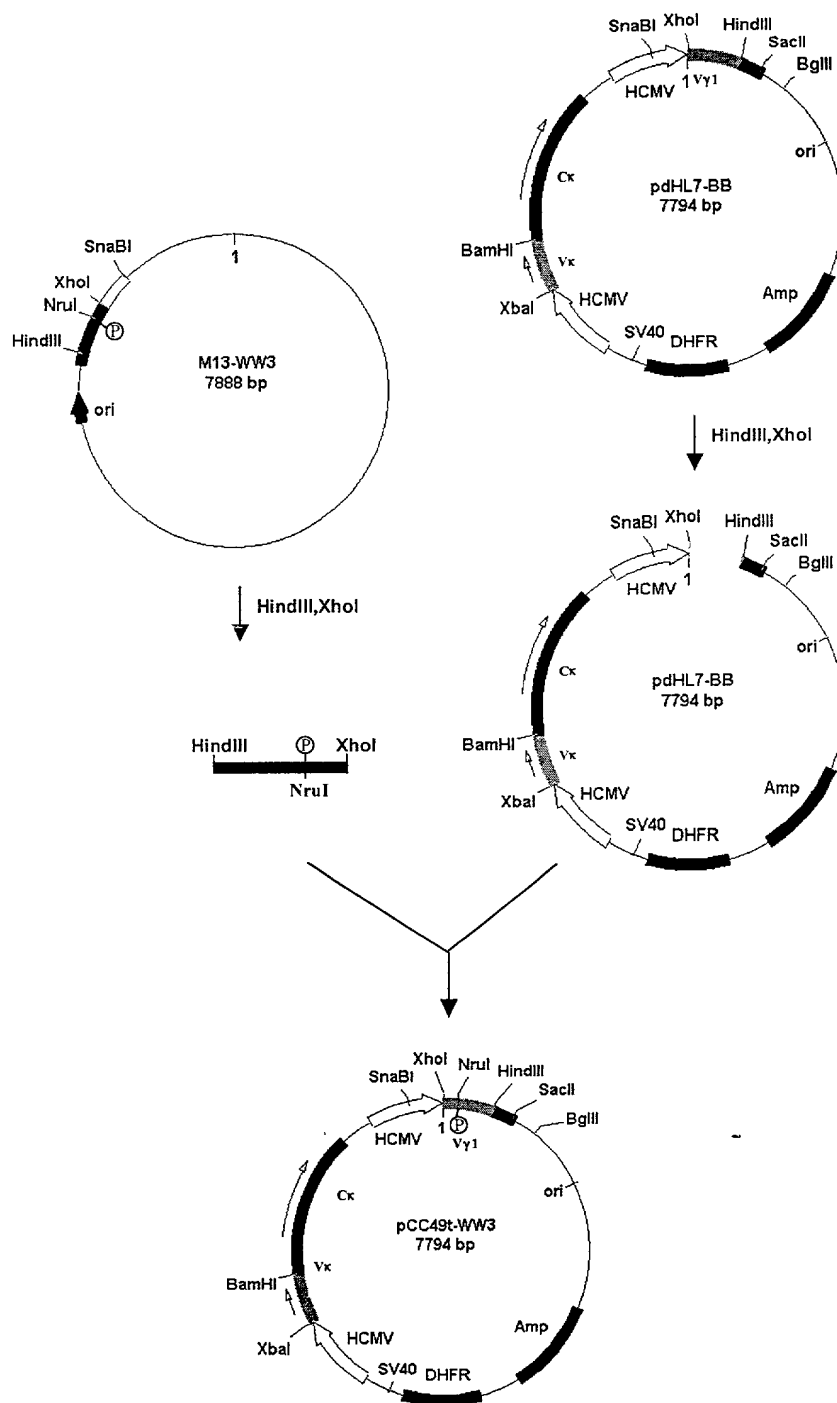


Fig. 20B

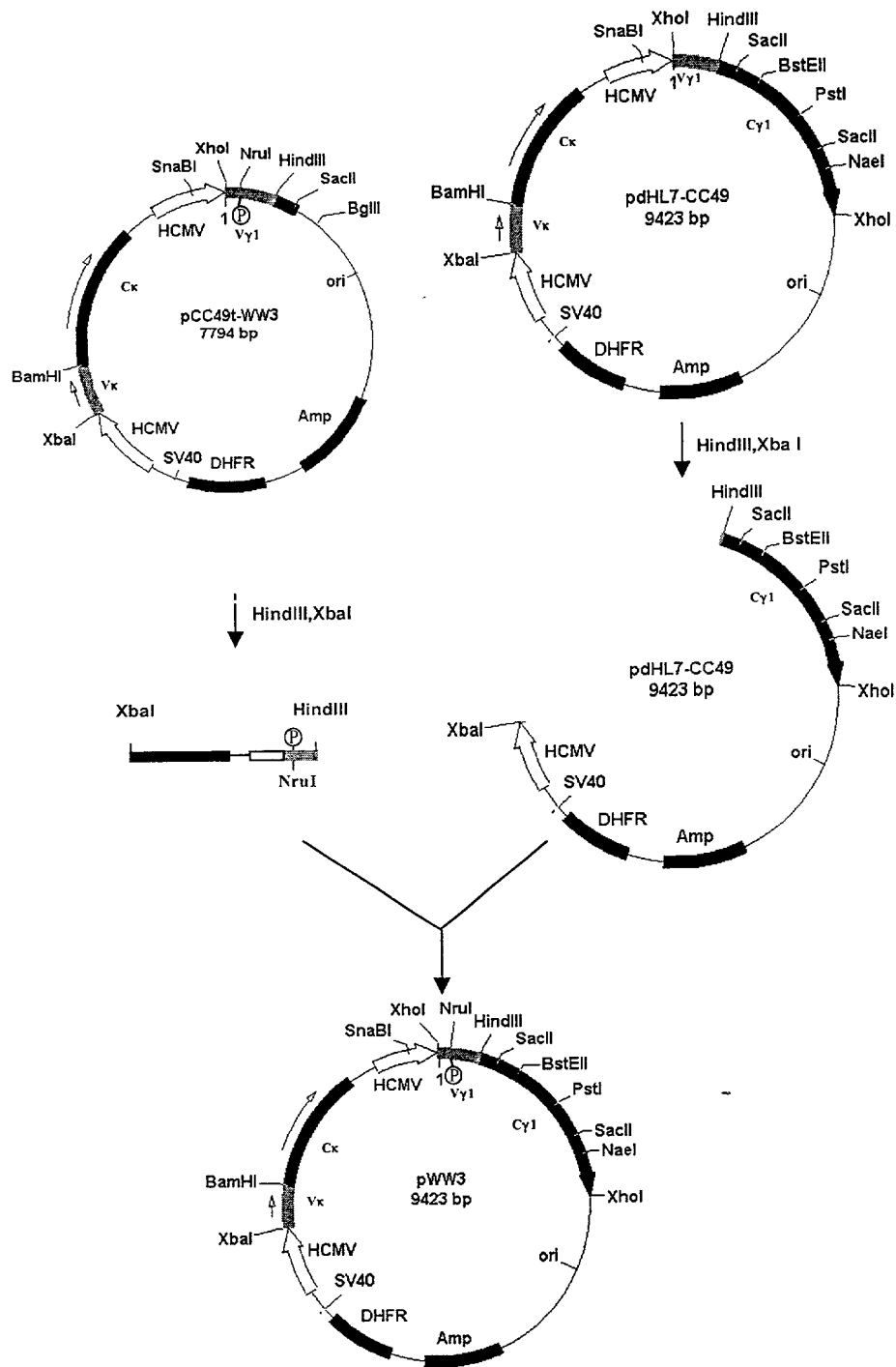


Fig. 20C

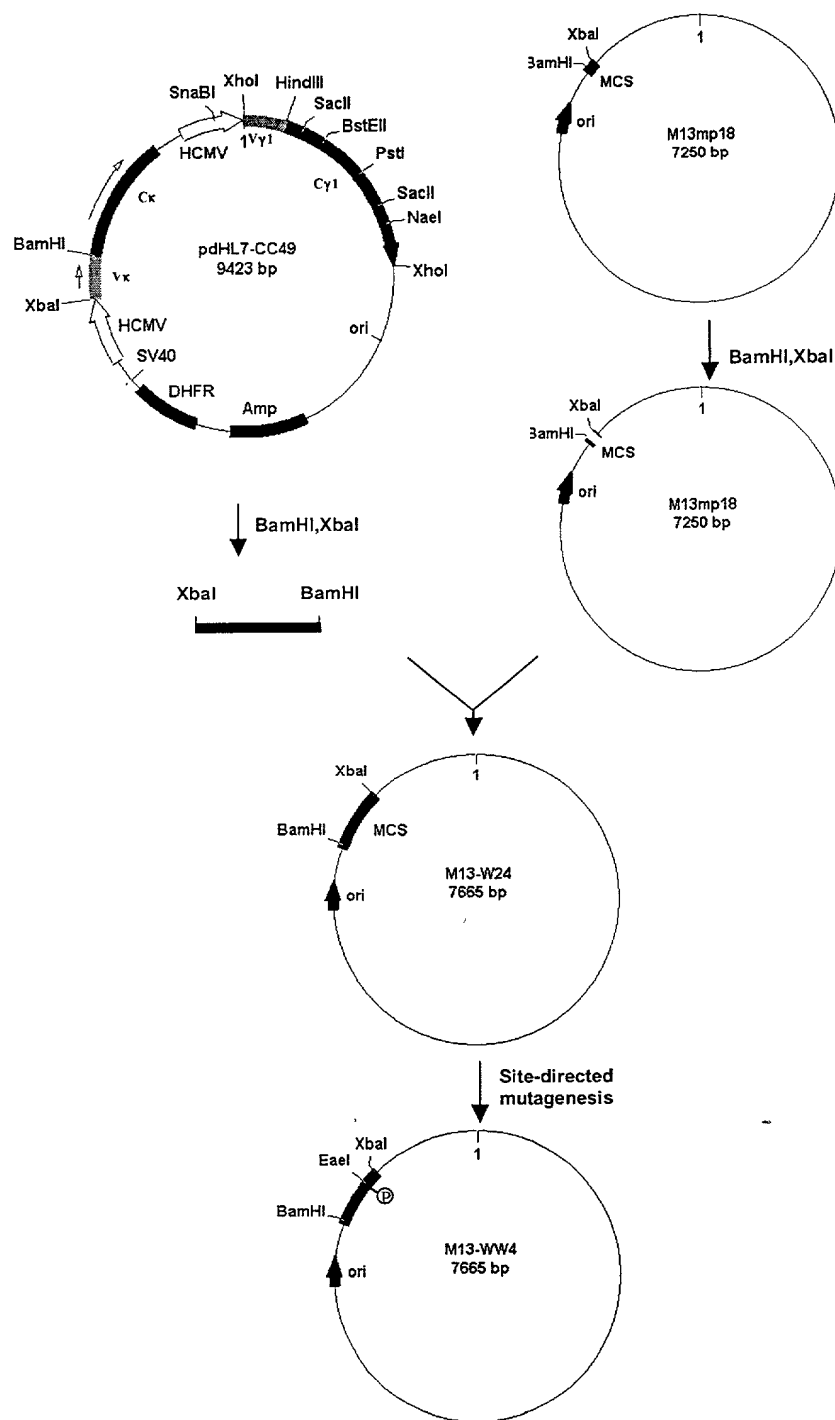


Fig. 21A

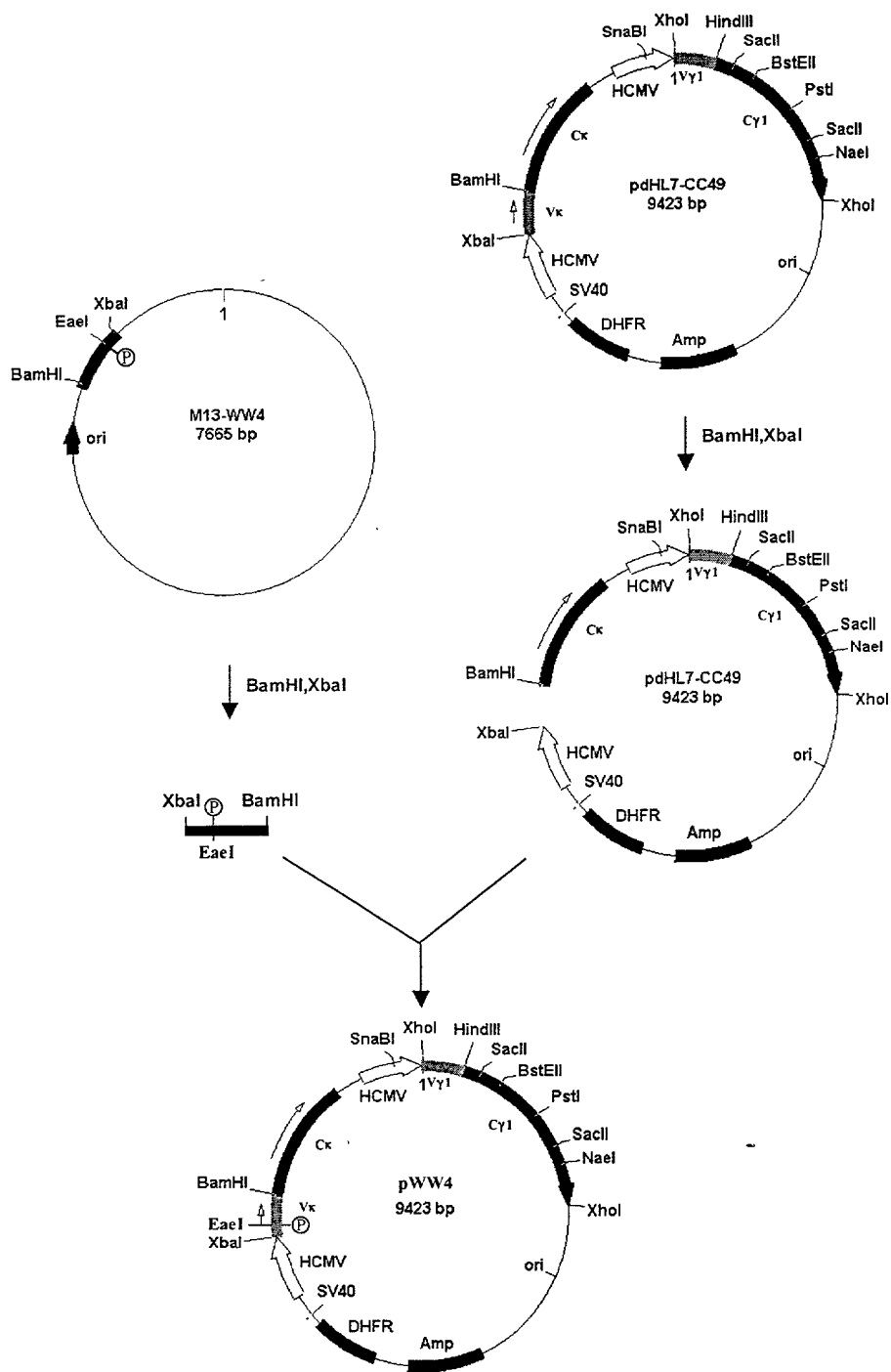


Fig. 21B

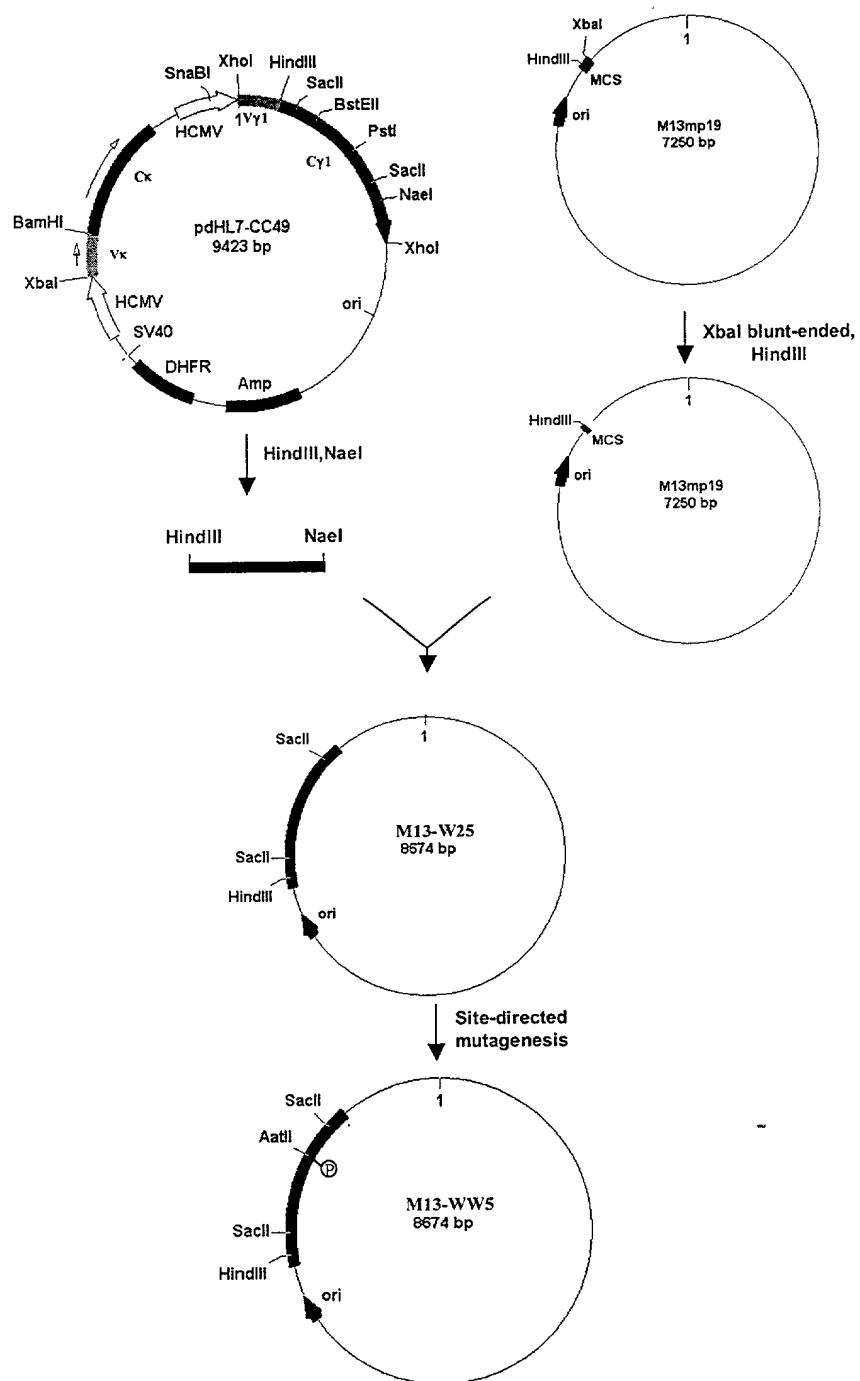


Fig. 22A

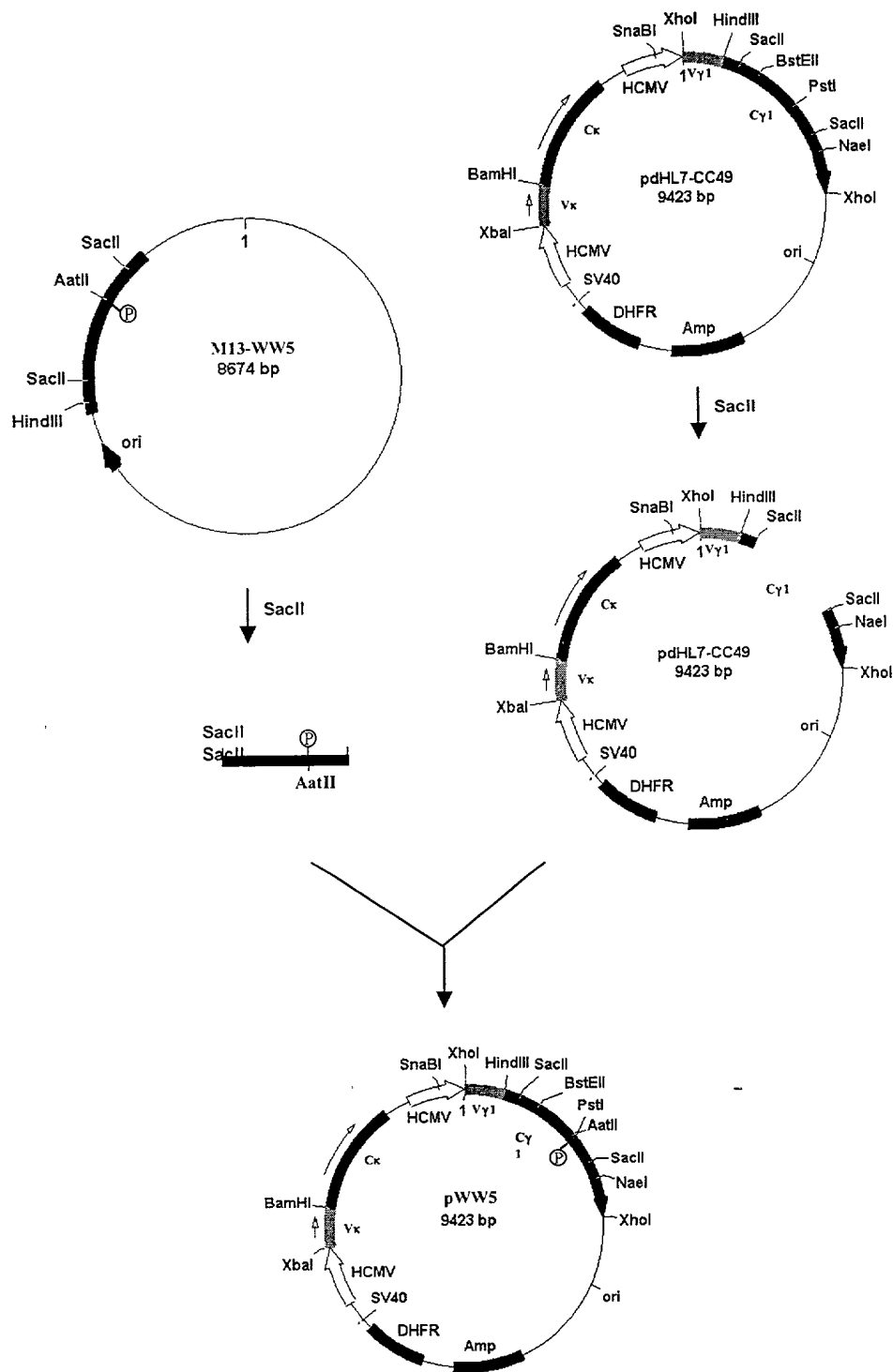


Fig. 22B

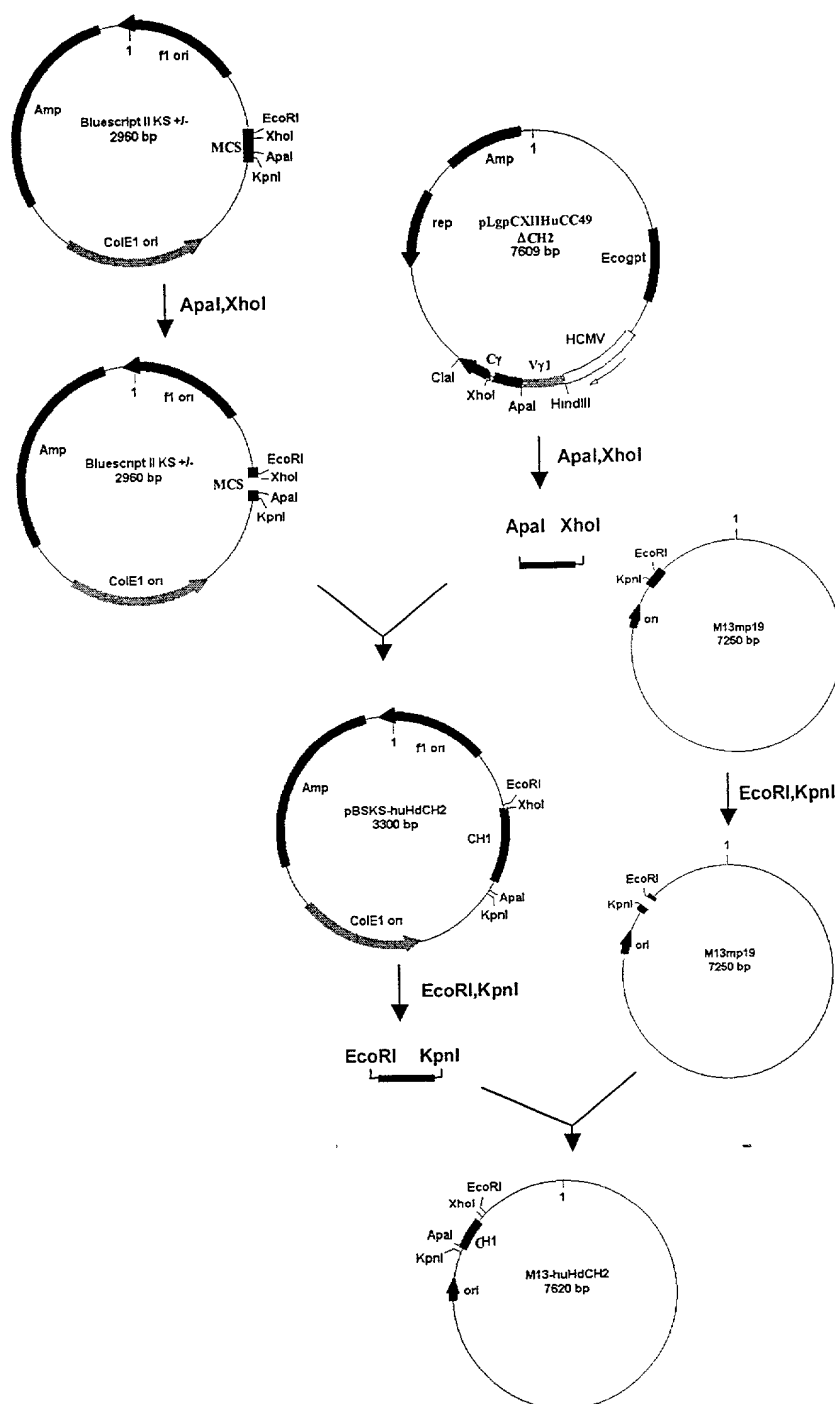


Fig. 23A

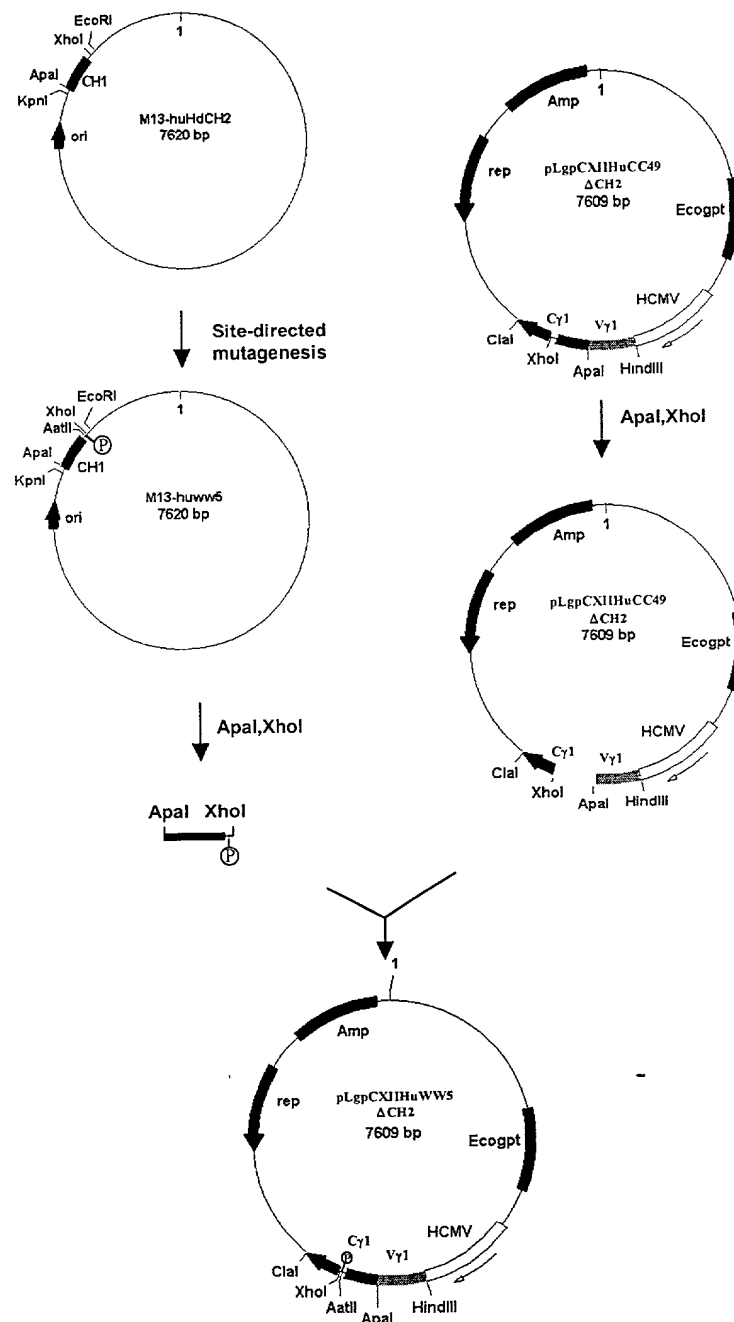


Fig. 23B

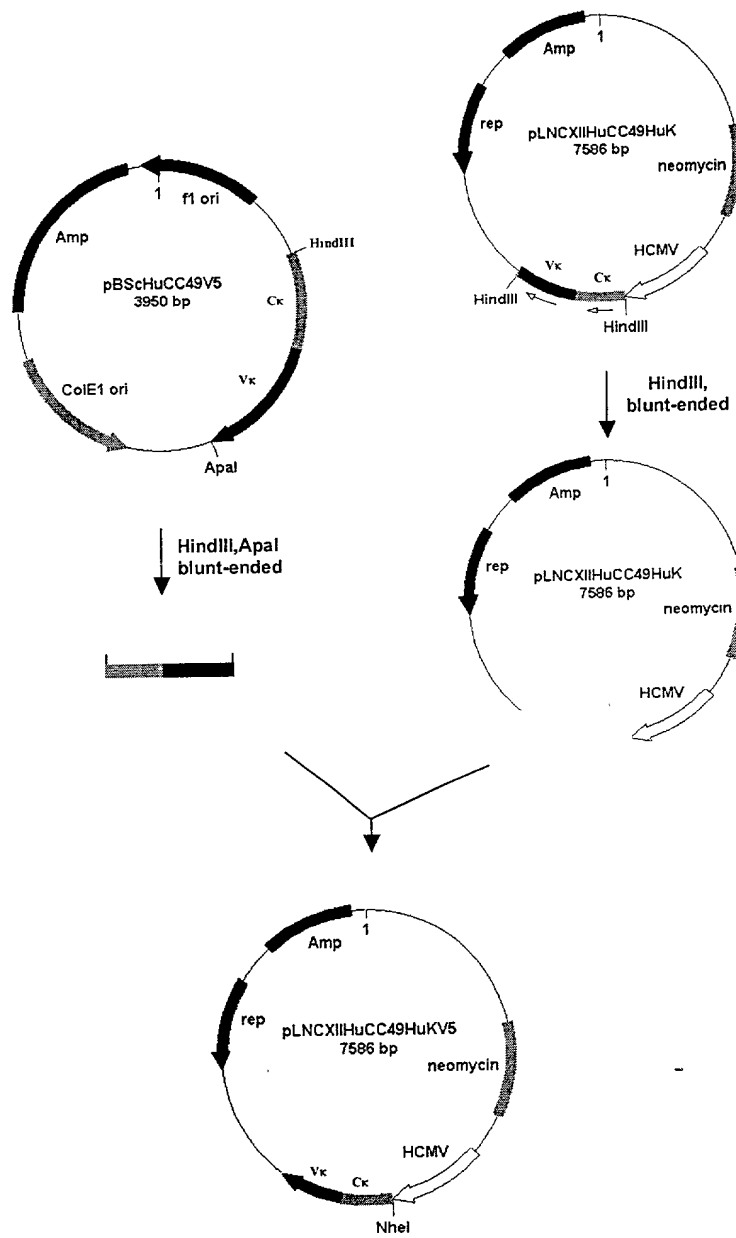


Fig. 24

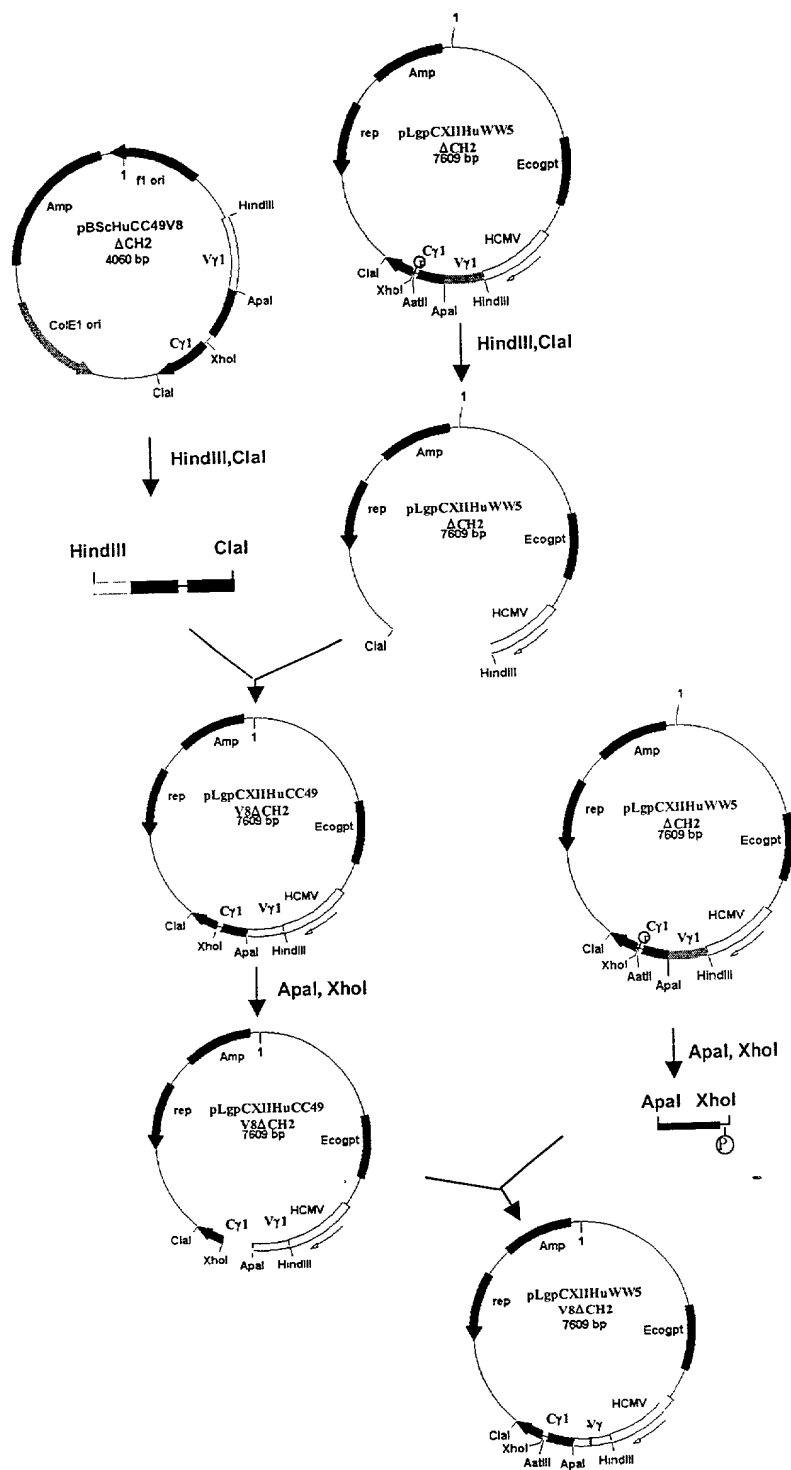


Fig. 25

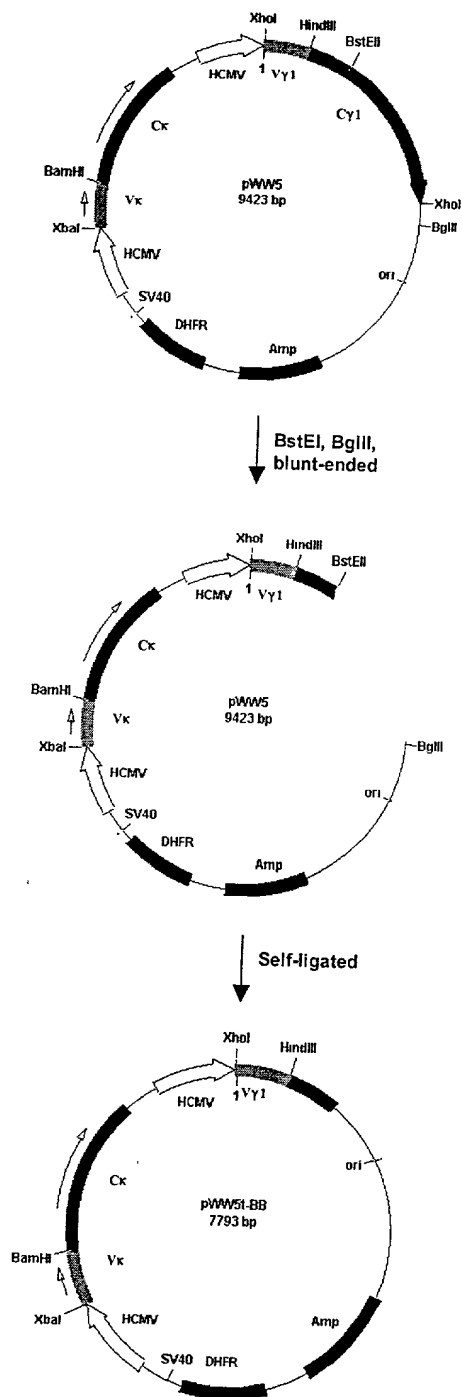


Fig. 26A

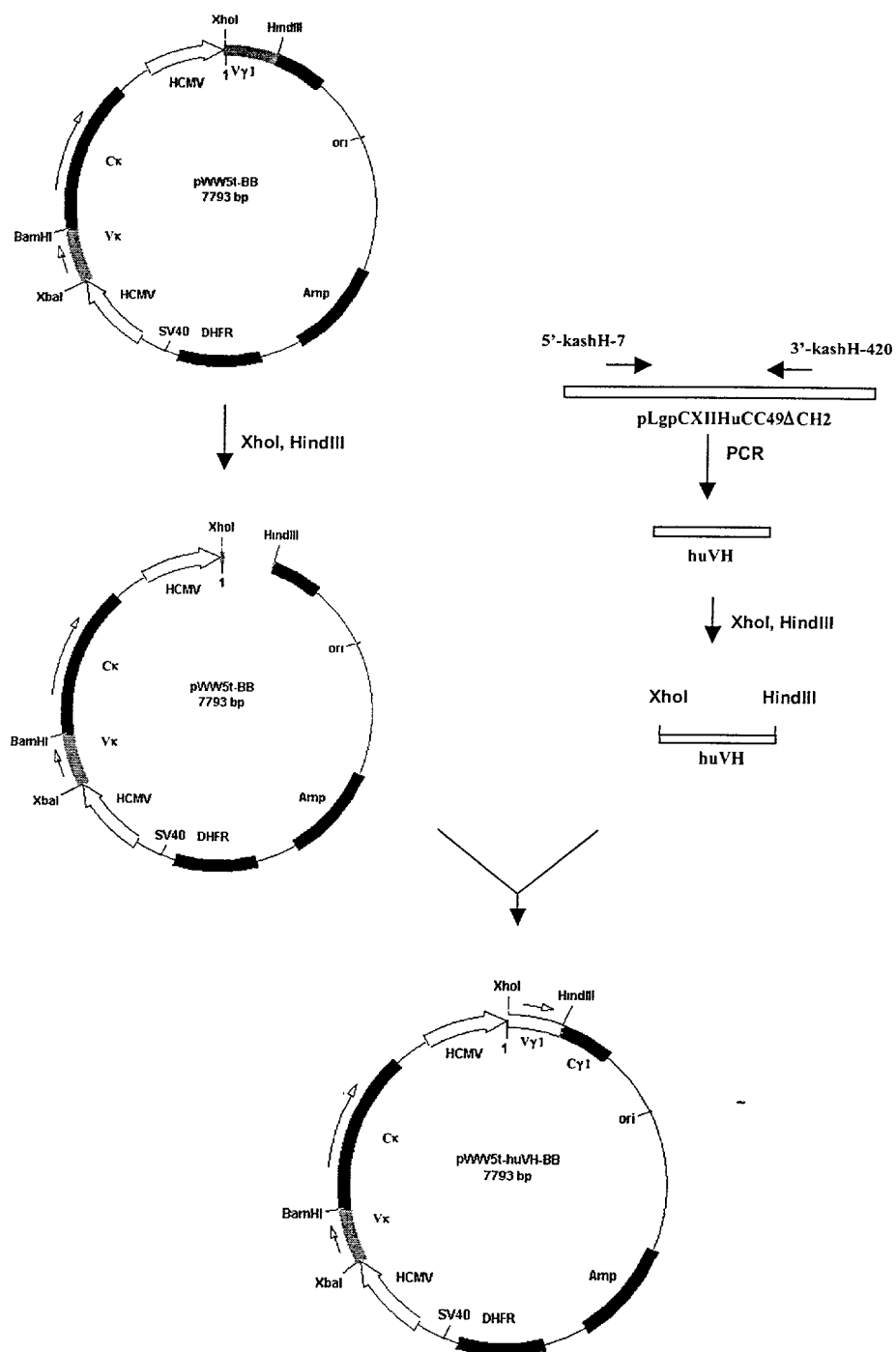


Fig. 26B

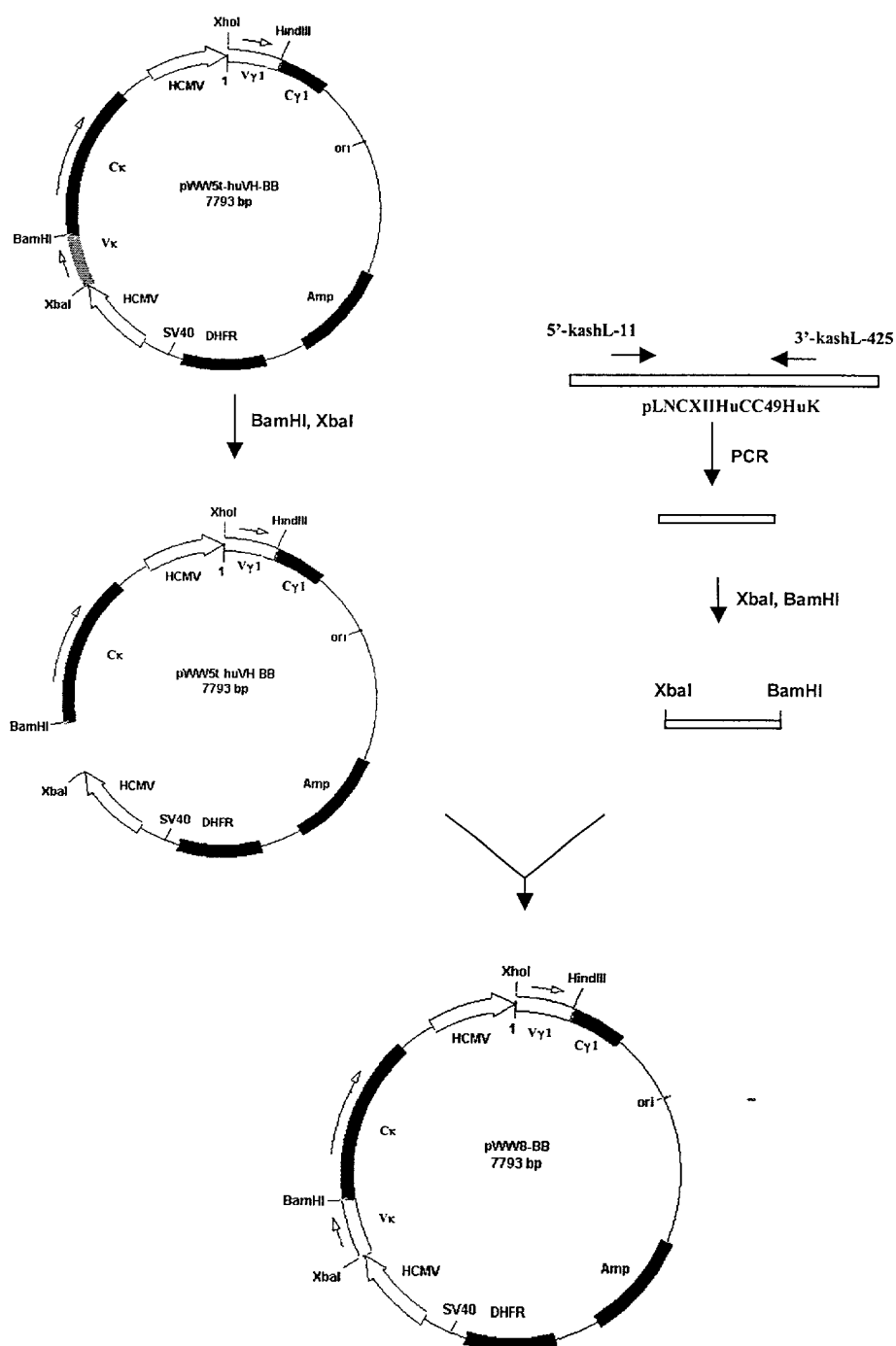


Fig. 26C

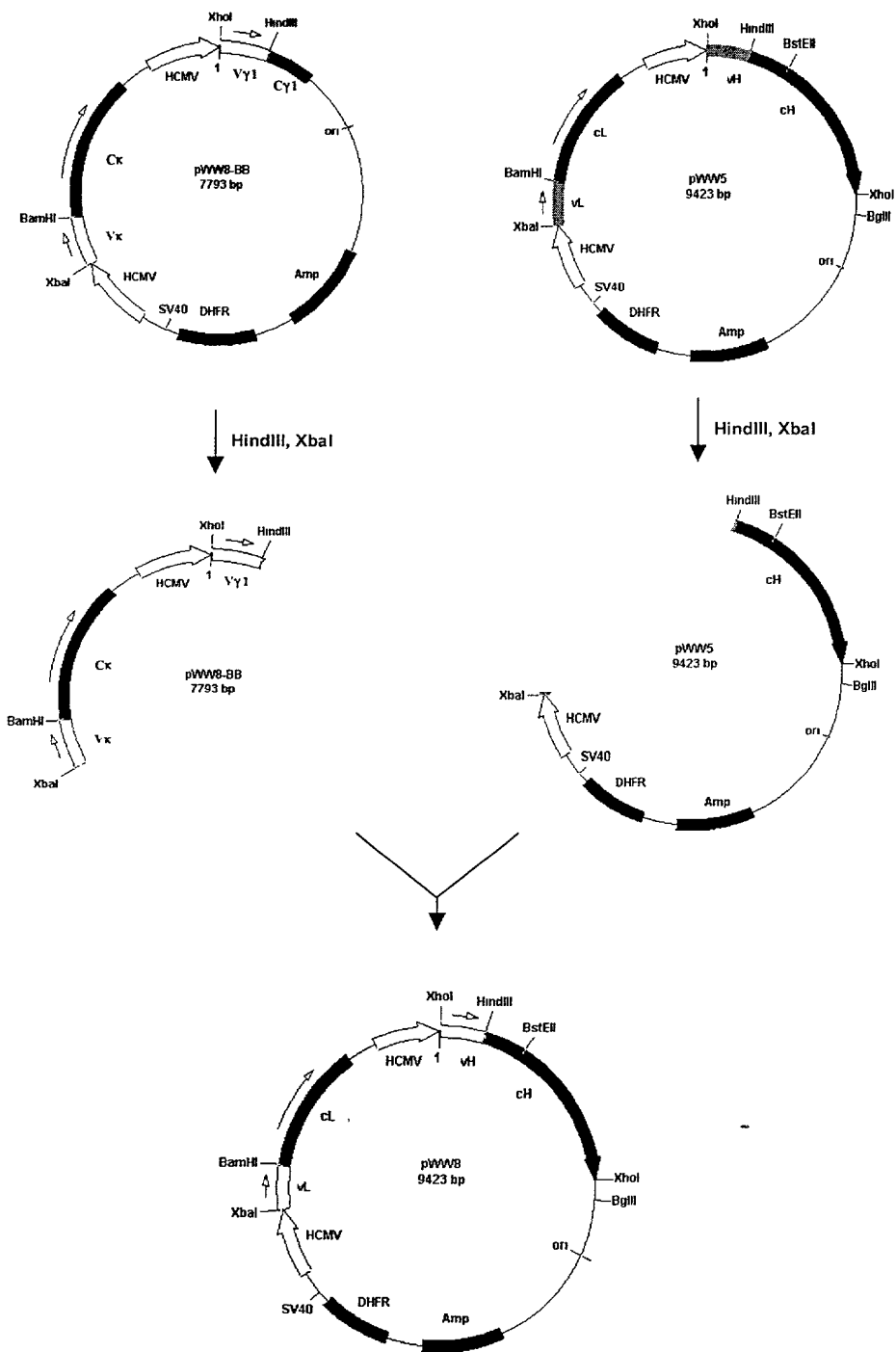


Fig. 26D

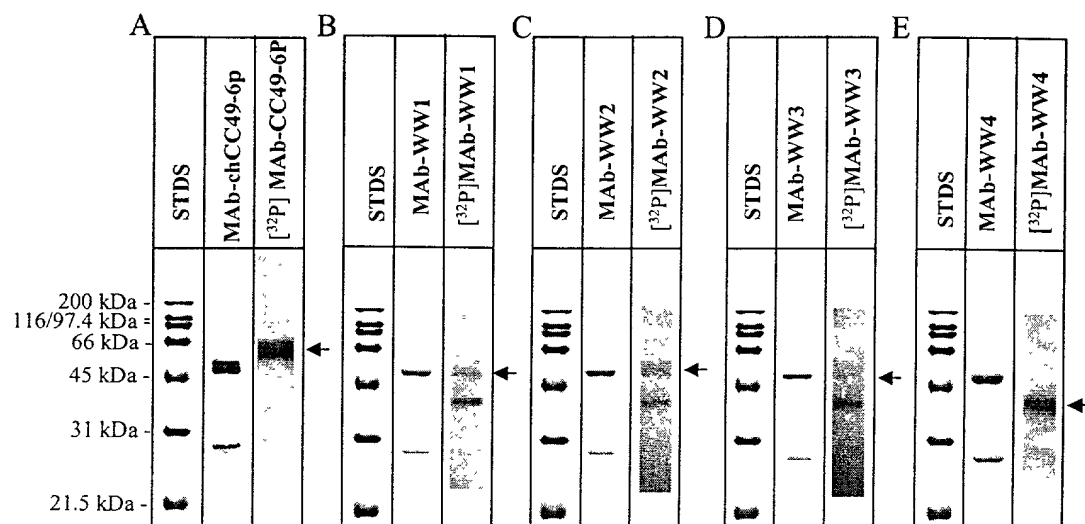


Fig. 27A-E

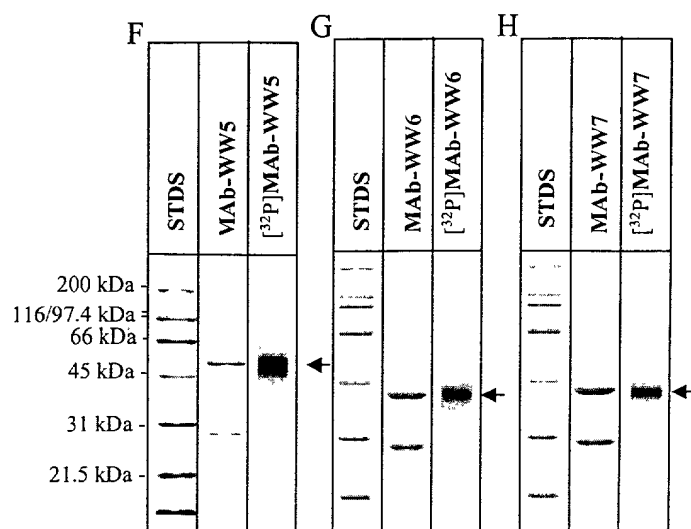


Fig. 27

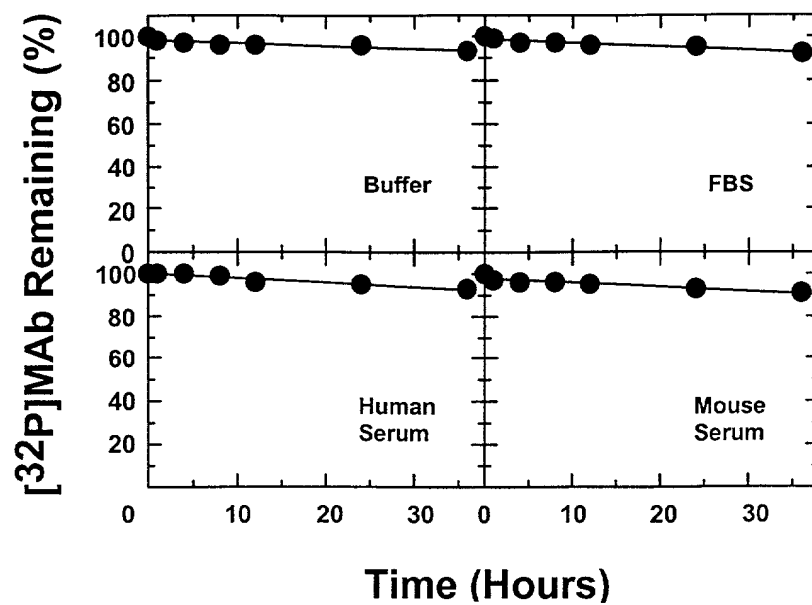


Fig. 28

TOP SECRET

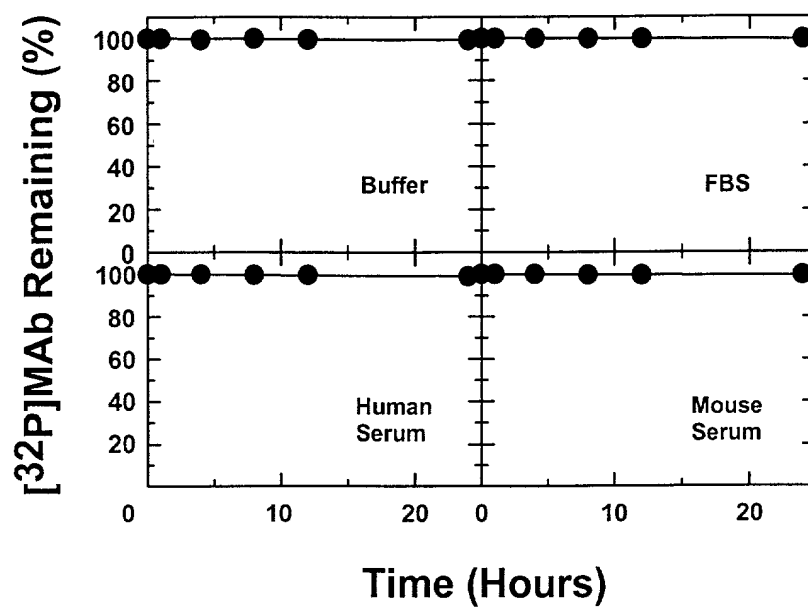


Fig. 29

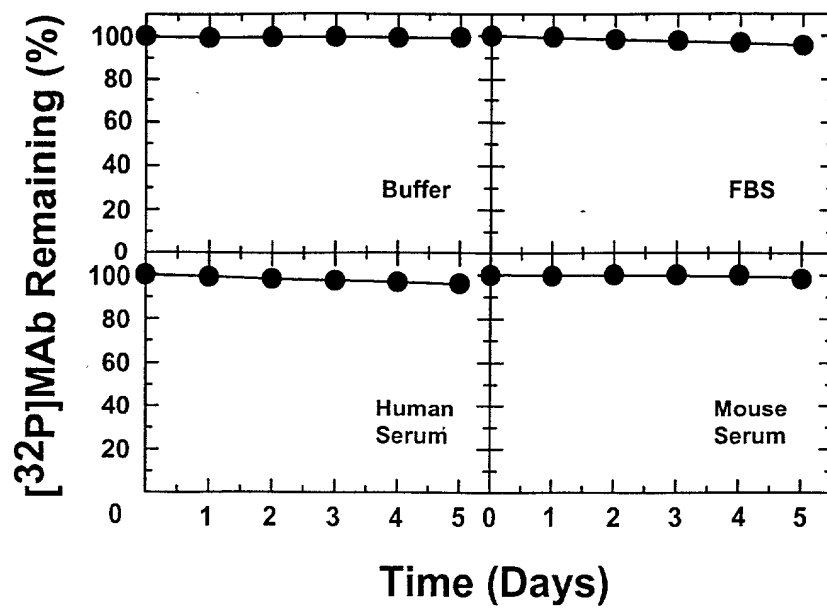


Fig. 30

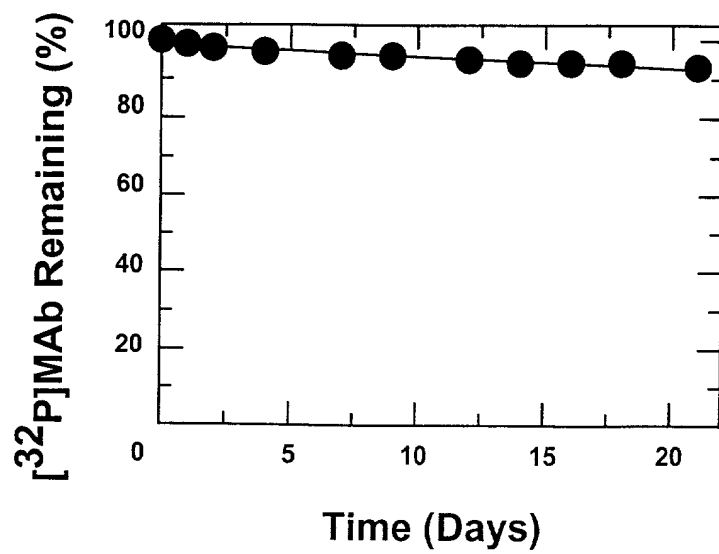


Fig. 31

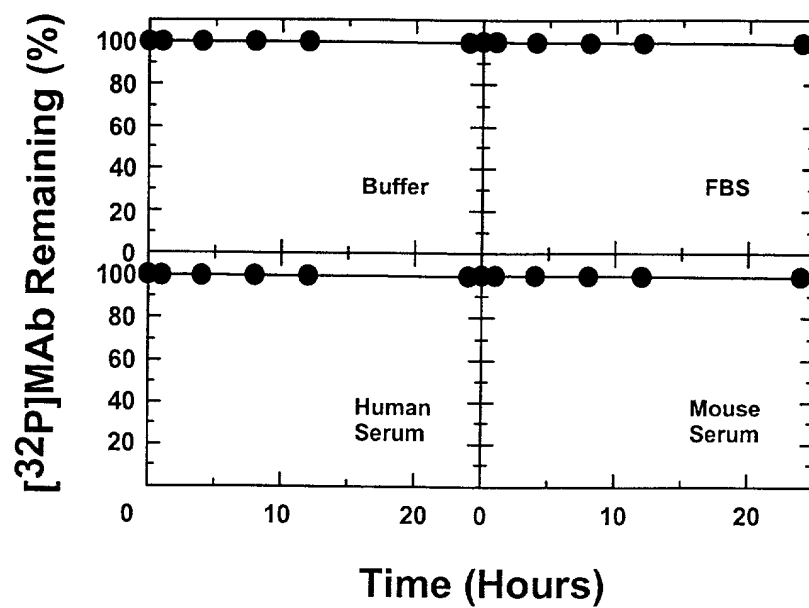


Fig. 32

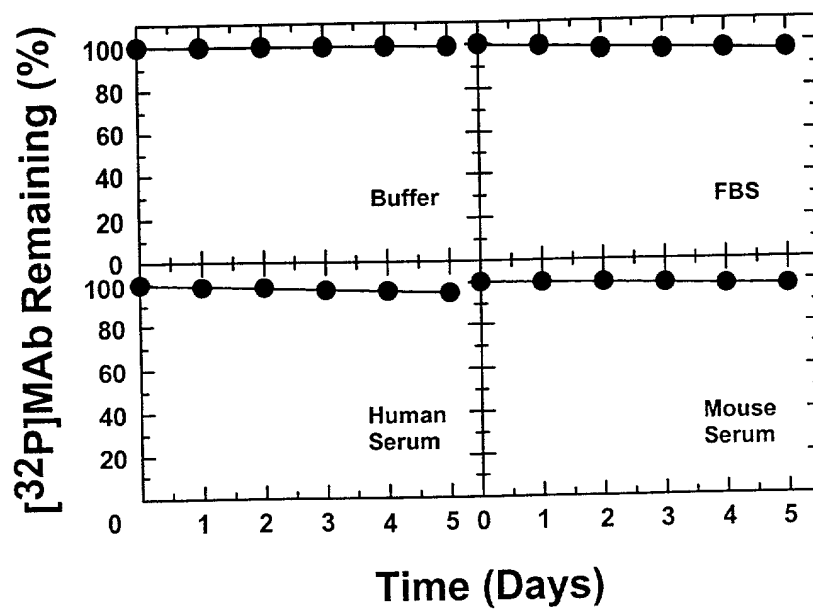


Fig. 33

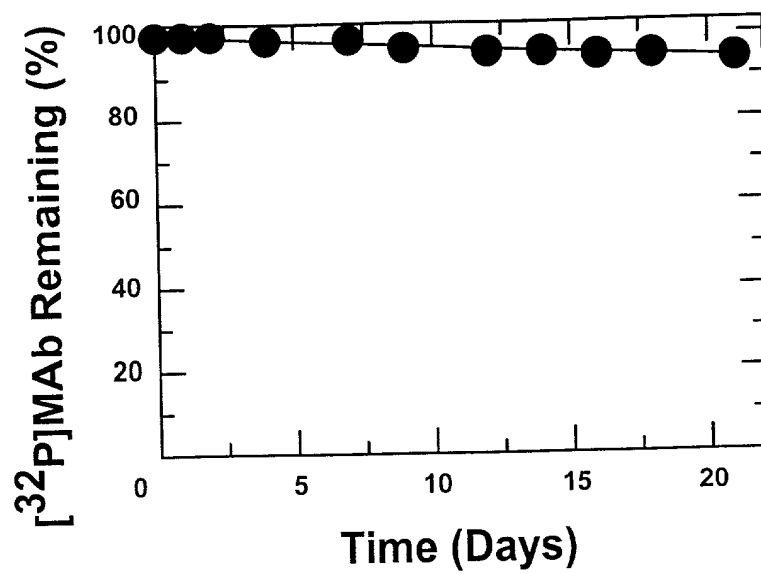
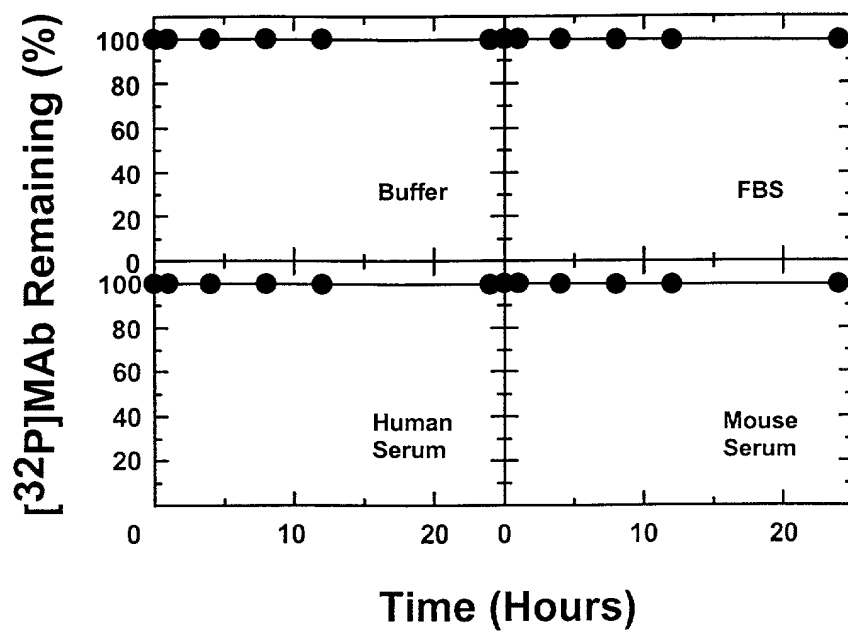


Fig. 34



P

Fig. 35

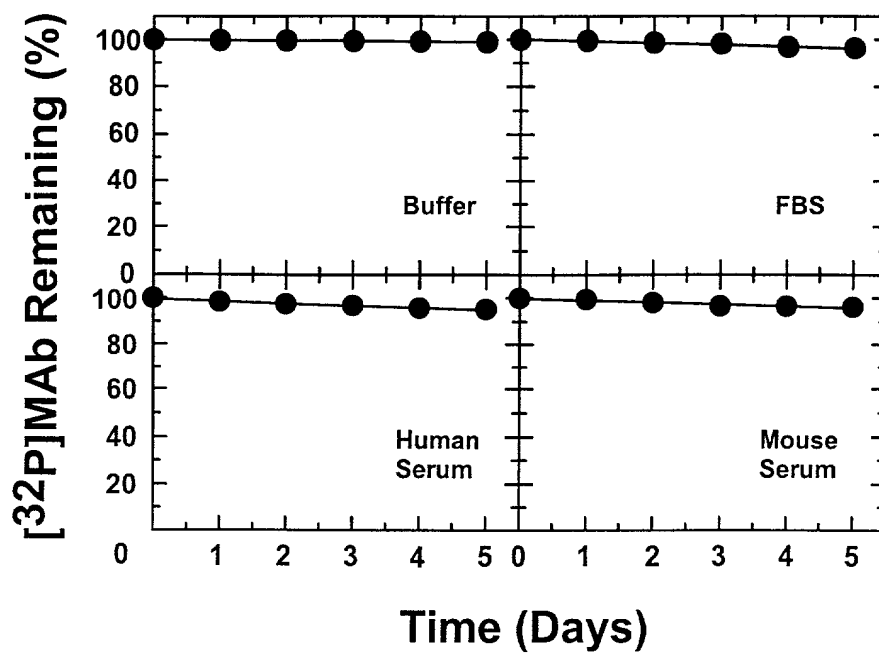


Fig. 36

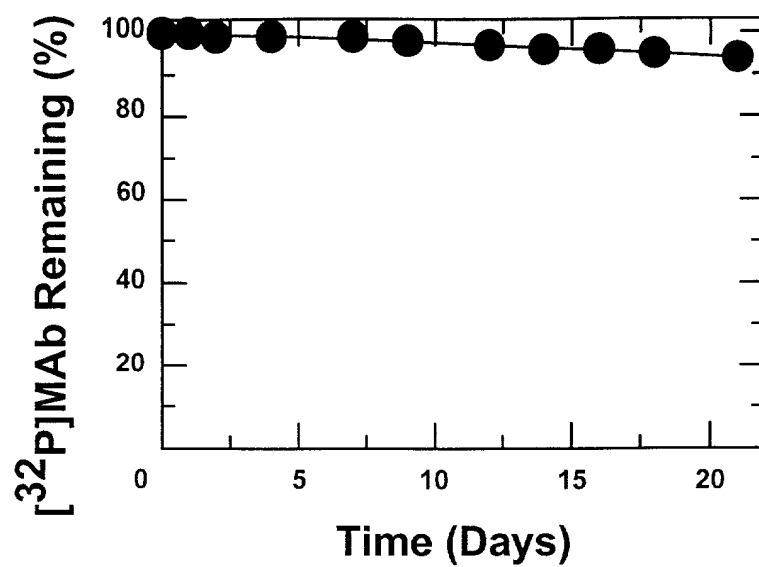


Fig. 37

093249-05101
T07E90-64E2E860

| | | Upper | Core | Lower |
|------------------------|-------------|-----------------|--------------|-------------|
| A: | MAb-chCC49: | EPKSCDKTHT | CPPCP | APELLGGP |
| | MAb231: | EPRGPTIKP | CPPCKCP | APNLLGGP |
| | MAb61.1.3: | VPRDCG | CKPCICT | VPEV |
| B: | | | | |
| MAb-chCC49 x MAb231 | | | | |
| | 1 | EPKSCDKTHTCPP | CP | APELLGGP 23 |
| | | | | |
| | 1 | EPRG | PTIKPCPPCKCP | APNLLGGP 24 |
| C: | | | | |
| MAb-chCC49 x MAb61.1.3 | | | | |
| | 1 | EPKSCDKTHTCPPCP | APELLGGP | 23 |
| | | | | |
| | 1 | VPRDCGCKPCICTV | VPEV | 17 |

Fig. 38

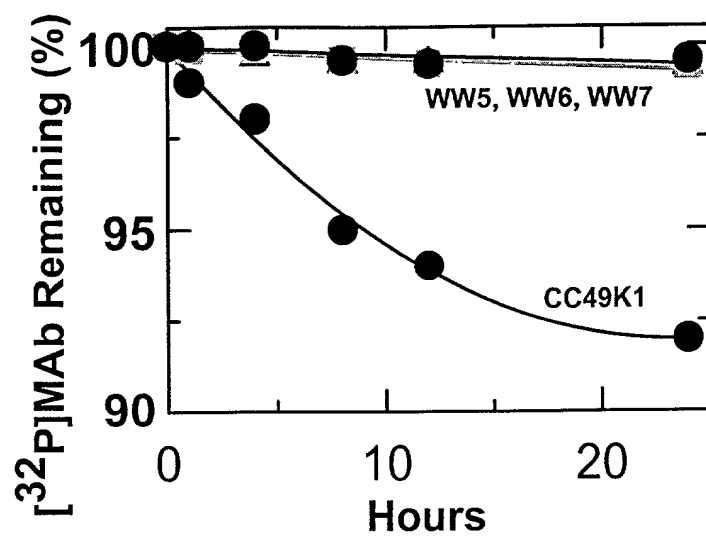


Fig. 39

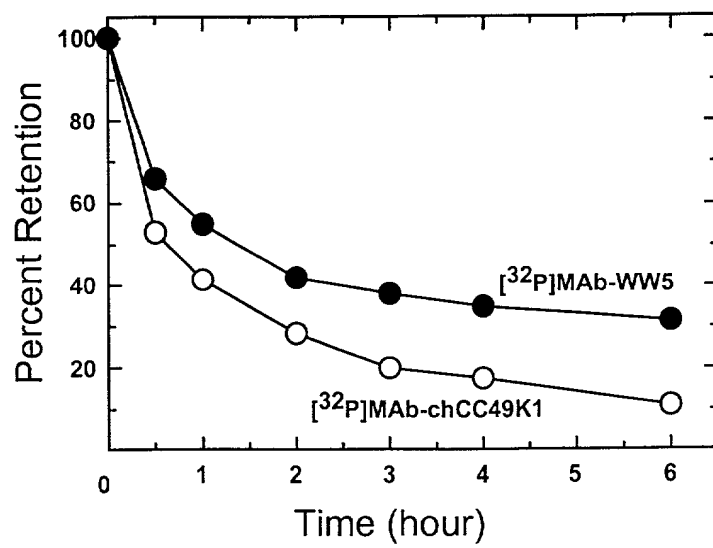


Fig. 40

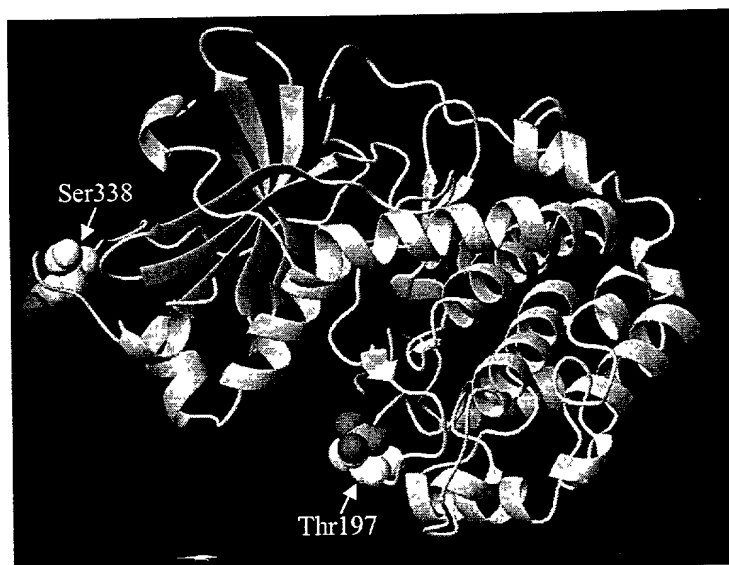


Fig. 41

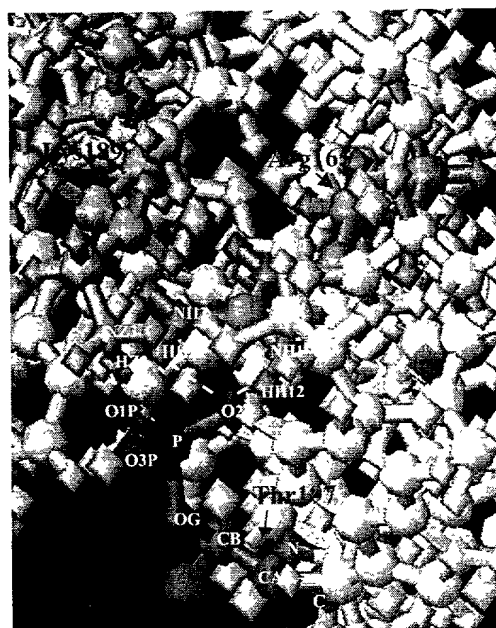


Fig. 42

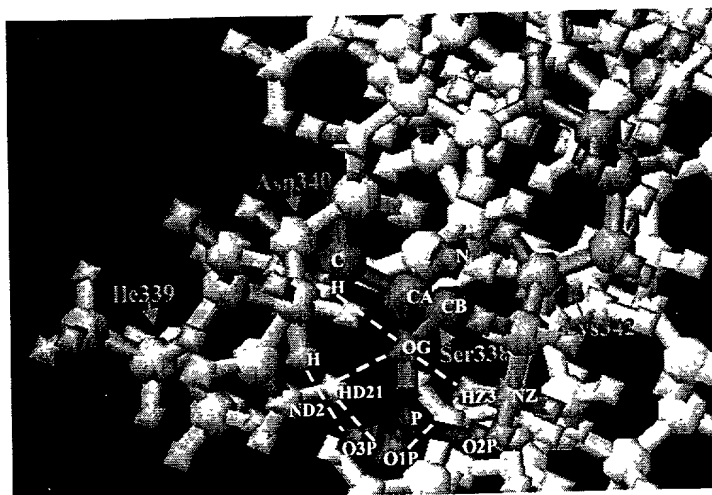


Fig. 43